SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Name:	nber 30	Examiner #:				
Art Unit: Phone Nun Mail Box and Bldg/Room Location:	nber 30Re	Serial Number: sults Format Preferred (circle):	PAPER DISK E-MAIL			
Mail Box and Bldg/Room Location: Results Format Preferred (circle): PAPER DISK E-MAII						
******	*****	******	*****			
Please provide a detailed statement of the sea Include the elected species or structures, key utility of the invention. Define any terms tha known. Please attach a copy of the cover shee	vords, synonyms, acr t may have a special	onyms, and registry numbers, and e meaning. Give examples or relevan	ombine with the concept or			
Title of Invention:	•					
Inventors (please provide full names):						
Earliest Priority Filing Date:						
*For Sequence Searches Only * Please include a appropriate serial number.			atent numbers) along with the			
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STAFF USE ONLY	Type of Search	Vendors and cost	where applicable			
Searcher: Devery C4994	NA Sequence (#)	STN	· · · · · · · · · · · · · · · · · · ·			
Searcher Phone #:	AA Sequence (#)	Dialog				
Searcher Location:	Structure (#)	Questel/Orbit				
Date Searcher Picked Up:	Bibliographic	Or.Link				
Date Completed: 08-08-9	Litigation	Lexis/Nexis				
Searcher Prep & Review Time:	Fulltext	Sequence Systems	· ·			
Clerical Prep Time:	Patent Family	CAN				
Online Time:	Other	Other (specify)				

PTO-1590 (1-2000)

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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648.2	1552 1261.6 198.6 790.2	Score
######################################	100.0 81.3 51.5 46.4	Query Match
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Sequence 37, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 5, Appli Sequence 4413, Ap Sequence 3, Appli Sequence 3, Appli Sequence 1, Appli	1 - 0 0 4 E	Description

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OTHER INFORMATION: Description of Artificial Sequence: Murine OTHER INFORMATION: surfactant protein D (without the CRD) fused to OTHER INFORMATION: the extracellular portion of human CD40L AUTHORS: Striggs, Melanie K.
AUTHORS: Striggs, Melanie K.
AUTHORS: Strockbine, L.
AUTHORS: Clifford, K.M.
AUTHORS: Clifford, K.M.
AUTHORS: Macduff, B.M.
AUTHORS: Macduff, B.M.
AUTHORS: Macduff, C.R.
TITLE: Recombinant human CD40 ligand stimulates B cell
JORNAL: J. Exp. Med.
TSGHF. K.
      ISSUE: 12
PAGES: 5671-5677
DATE: 1995
RELEVANT RESIDUES: 32 TO 800
US-09-454-223-1
                                                                              JOURNAL: J.
                                                                                                  RELEVANT RESIDUES: 801 TO 1600
PUBLICATION INFORMATION:
AUTHORS: MOTWANT, M
TITLE: Mouse surfactant protein-D.
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US-09-454-223-1
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TITLE OF INVENTION: Multimeric forms of CD40L and other TNF family members
FILE REFERENCE: TNFSF-collectin fusion proteins
CURRENT APPLICATION NUMBER: US/09/454,223
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEY: CDS
LOCATION: (32)..(1444)
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LOCATION: (7)..(31)
FEATURE:
NAME/KEY: misc_feature
LOCATION: (88)..(799)
OTHER INFORMATION: Mature murine surfactant protein D including hub
OTHER INFORMATION: region, collagenous portion, and neck, but
OTHER INFORMATION: excluding carbohydrate recognition domain (CRD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PEATURE:
NAME/KEY: sig_peptide
LOCATION: (32)...(88)
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GENERAL INFORMATION:
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LOCATION: (801)..(1546)
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      NAME/KEY: misc_recomb
LOCATION: (88)..(799)
OTHER INFORMATION: Matu
OTHER INFORMATION: redd
OTHER INFORMATION: excl
                                                                                                                                                   FEATURE:

NAME/KEY: 81g_peptide
LOCATION: (32)..(88)
OTHER INFORMATION: S1gr
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                                                                                                                                                                                                                                                                                  OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                   NAME/KEY: 5'UTR
LOCATION: (7)..(31)
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
                                                                                                       NAME/KEY: CDS
LOCATION: (32
                                                                                        FEATURE:
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ION: (88)...(799)
INFORMATION: Mature murine surfactant protein D including INFORMATION: region, collagenous portion, and neck, but INFORMATION: excluding carbohydrate recognition domain (CR
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                                                                                                       (32)..(1441)
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                                                                                                                                                                                                                                                                                  Description of Artificial Sequence: Murine surfactant protein D (except CRD) fused to extracellular domain of murine CD40 ligand
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JOURNAL: Nature
VOLUME: 357
ISSUE: 6373
PAGES: 80-82
DATE: 1992
RELEVANT RESIDUES: 8
US-09-454-223-5
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Best Local Similarity
Matches 1349; Conserv
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AUTHORS: White, R A.
AUTHORS: Guo, N
AUTHORS: Dowler, L L.
AUTHORS: Tauber, A I.
AUTHORS: Motwani, M
TITLE: characterization, a
TITLE: 14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature LOCATION: (800)..(1441) OTHER INFORMATION: Murine OTHER INFORMATION: stalk PUBLICATION INFORMATION:
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PUBLICATION INFORMATION:
AUTHORS: Armitage, R
TITLE: Molecular and bio
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VOLUME: 155
ISSUE: 12
PAGES: 5671-5677
DATE: 1995
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RESULT 3
US-09-454-223-3
Sequence 3, Application US/09454223
GENERAL INFORMATION:
APPLICANT: ROZOBLUTH, Richard S
TITLE OF INVENTION: Multimeric forms of CD40L and other
FILE REFERENCE: TNFSF-collectin fusion proteins
CURRENT APPLICATION NUMBER: US/09/454,223
CURRENT FILING DATE: 1999-12-09
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LOCATION: (801)..(1534)
OTHER INFORMATION: Murine RANKL/TRANCE extracellular region,
OTHER INFORMATION: including stalk
OTHER INFORMATION:
OTHER INFORMATION:
AUTHORS: MOLWANI, M
AUTHORS: MOLWANI, M
AUTHORS: Dowler, L L.
AUTHORS: Dowler, L L.
AUTHORS: Tauber, A I.
AUTHORS: Motwani, M
TITLE: Mouse surfactant protein-D. cDNA cloning,
TITLE: id.
TOTHERS: 14.
TO
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Best Local (
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TITLE: A homologue of the TNF receptor and its ligand
JOURNAL: Nature
VOLUME: 390
ISSUE: 6656
PAGES: 175-179
DATE: 1997
DATE: 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PUBLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ISSUE: 12
PAGES: 5671-5677
DATE: 1995
                     ORGANISH: Artificial Sequence FEATURE: NAME/KEY: 5'UTR LOCATION: (7)..(31) OTHER INFORMATION: 5' UTR taker OTHER INFORMATION: protein D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JOURNAL: J. Immunol.
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SOFTWARE: PatentIn Ver.
SEQ ID NO 3
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NAME/KEY: misc_feature
LOCATION: (32)..(800)
OTHER INFORMATION: Murine surfactant protein D including hub region,
OTHER INFORMATION: collagenous portion, and neck, but excluding
OTHER INFORMATION: carbohydrate recognition domain (CRD)
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LOCATION: (32)..(88)
OTHER INFORMATION: Sig
                                                                                                                                                                                                                            LOCATION: (32)..(1534)
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NAME/KEY: CDS
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Pred. No. 1.5e-185;
0; Mismatches 9;
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APPLICANT: AHUJA, SEEMA
APPLICANT: BONEWALD, LYNDA
TITLE OF INVENTION: CD40 LIGAND AND CD40 AGONIST COMPOSITIONS AND METHODS OF US
FILE REFERENCE: 4003.001000
CURRENT APPLICATION NUMBER: US/09/645,926A
CURRENT FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: 60/151,250
PRIOR APPLICATION NUMBER: 50/151,250
PRIOR FILING DATE: 1999-08-27
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.0
SEQ ID NO 1
LENGTH: 879
TYPE: DNA
ORGANISM: Homo sapiens
US-09-645-926A-1
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                                             Query Match
Best Local Similarity
Matches 722; Conserv
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                                              Conservative
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                                                         Score 720.2; DB 25
Pred. No. 2.3e-166;
                                              Mismatches
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                                                                    SOFTWARE: P
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         TYPE: DNA
ORGANISM: HOMO
FEATURE:
NAME/KEY: CDS
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                                                          LENGTH:
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APPLICANT: L1u, Lina
APPLICANT: Lotze, Michael T
ITTLE OF INVENTION: GENETICALLY-MODIFIED ANTIGEN PRE
TITLE OF INVENTION: INDUCTION OF IMMUNOTOLERANCE AND
FILE REFERENCE: UPT-003
CURRENT APPLICATION NUMBER: US/09/469,519
CURRENT FILING DATE: 1999-12-22
EARLIER APPLICATION NUMBER: 60/113766
EARLIER APPLICATION NUMBER: -60/113766
EARLIER FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 38
NUMBER OF SEQ ID NOS: 38
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LOCATION: (112)..(183)
OTHER INFORMATION: NH2 signal/ anchor
TITLE: Recombinant human CD40 ligand stimulates B cell
TUBLICATION INFORMATION:
TITLE: Proliferation and immunoglobulin E secretion
VOLUME: 176
ISSUE: 6
PAGES: 1543-1550
DATE: 1992
DATABASE ACCESSION NUMBER: X67878; S50586
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Best Local Similarity
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98.6%;
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Pred. No. 9.5e-164;

 Mismatches

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PRIOR APPLICATION NUMBER: 09/816
PRIOR FILING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 584
SOFTWARE: Custom
SEQ ID NO 373
LENGTH: 1816
TYPE: DNA
ORGANISM: Homo Sapiens
PCT-US01-08655-373
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CURRENT FILING DATE: 2001-04-16
PRIOR APPLICATION NUMBER: 09/522,929
PRIOR FILING DATE: 2000-04-18
PRIOR APPLICATION NUMBER: 09/770,160
PRIOR FILING DATE: 2001-01-26
PRIOR FILING DATE: 2001-01-26
PRIOR FILING DATE: 2000-09-22
PRIOR FILING DATE: 2000-09-22
PRIOR FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: 09/695,783
PRIOR FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: 09/728,628
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2001-02-1783,066
PRIOR FILING DATE: 2001-02-1783,066
PRIOR APPLICATION NUMBER: 09/78,026
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Best Local Similarity
Matches 716; Conserv
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PCT-US01-08655-373
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                                                 actacaccatgagcaacaacttggtaaccctggaaaatgggaaacagctgaccgttaaaa
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Pred. No. 9.5e-164;
0; Mismatches 10;
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CURRENT APPLICATION NUMBER: PCT/US99/30930
CURRENT FILING DATE: 1999-12-22
EARLIER APPLICATION NUMBER: U.S. 60/114,106
EARLIER FILING DATE: 1998-12-29
NUMBER OF SEQ ID NOS: 2
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 1816
TYPE: DNA
ORGANISM: Homo Sapiens
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PCT-US99-30930-1
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Best Local Similarity 98.6%;
Matches 716; Conservative
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Pred. No. 9.5e-164;
0; Mismatches 10;
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; LENGTH: 1816
; TYPE: DNA
; ORGANISM: HOMO !
US-09-470-494-1
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APPLICANT: Wagner, David H.
APPLICANT: Newell, Evan
TITLE OF INVENTION: Use of CD40 Engagement to Al
TITLE OF INVENTION: Receptor Usage
FILE REFERENCE: IO277/7007/HCL/KA
CURRENT APPLICATION NUMBER: US/09/470,494
CURRENT FILING DATE: 1999-12-22
EARLIER APPLICATION NUMBER: U.S. 60/114,106
EARLIER FILING DATE: 1998-12-29
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US-09-470-494-1
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Best Local Similarity
Matches 716; Conserv
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SOFTWARE: FastSEQ for
SEQ ID NO 1
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Pred. No. 9.5e-164;
Pred. No. 9.5e-164;
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APPLICANT: Wagner, David H.

APPLICANT: Newell, Evan

ITILE OF INVENTION: Use of CD40 Engagement to

ITILE OF INVENTION: Receptor Usage

CURRENT APPLICATION NUMBER: US/09/470,494A

PRIOR APPLICATION NUMBER: US/09/470,494A

PRIOR APPLICATION NUMBER: US. 60/114,106

PRIOR FILING DATE: 1999-12-29

PRIOR FILING DATE: 1998-12-29

SOPTWARE: FastSEQ for Windows Version 3.0

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TYPE: DNA
ORGANISM: Homo Sapiens
US-09-470-494A-1
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Best Local Sim
Matches 716;
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US-09-470-494A-1
VS-09-470-494A-1
Sequence 1, Application US/09470494A
GENERAL INFORMATION:
GENERAL INFORMATION:
Martha Karen
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                            293 aggagattaaaagccagtttgaaggctttgtgaaggatataatgttaaacaaagaggaga
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                                                                                                                                                                                                       ilarity 98.6%;
Conservative
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Pred. No. 9.5e-164; Length 1816;
0; Mismatches 10; Indels 0
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TYPE: DNA
ORGANISM: Homo sapiens
US-09-645-926A-5
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: APULA, SEEMA
APPLICANT: APULATION:
CD40 LIGAND AND CD40 AGONIST COMPOSITIONS AND METHODS OF (
TITLE OF INVENTION:
CURRENT APPLICATION UNMBER: US/09/645,926A
CURRENT APPLICATION NUMBER: 60/151,250
PRIOR APPLICATION NUMBER: 60/151,250
NUMBER OF SEQ ID NOS: 7
SOFTWARE:
SEQ ID NO 5
LEMCTH: 1816
                                                                                                                          Query Match
Best Local Sim
Matches 716;
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                                                 173 alottoatagaaggitggacaagatagaagatgaaaggaalottoatgaagatittgiat 232
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                                                                                                              Score 710; DB 25;
Pred. No. 9.5e-164;
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CURRENT APPLICATION NUMBER: US/09/577,408

CURRENT FILING DATE: 2000-05-18

NUMBER OF SEQ ID NOS: 8502

SOFTWARE: PL_GCT_genes Version 1.0

SEQ ID NO 4413

LENGTH: 1833

TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1388)...(1709)
OTHER INFORMATION: Similar to gill10587 in the genepept database relations of the control 
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APPLICANT: Sinku, Ankura
APPLICANT: Liu, Chenghua
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: Novel Nucleic Ac
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 792
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COMPUTER

READABLE FORM

ADDRESSEE: STERN,
STREET: 100 NEW Y
CITY: WASHINGTON
STATE: D.C.
SIP: 20005

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RESULT 12
US-08-234-580-3
; Sequence 3, Application US/08234580
; GENERAL INFORMATION:
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Best Local S
Matches 713
APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING
TITLE OF INVENTION: DIFFERENTIATING B CELLS, A
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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98.2%;
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Pred. No. 1.4
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LOCATION:
US-08-234-580-3
1395 gccaagtgagccatggcactggcttcacgtcctttggcttactcaaactctgaacagtgt 1454
                                                                         Query Match

Best Local Similarity 99.6%; Score 681.2; DB 6;
Matches 683; Conservative 0; Mismatches 3;
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, V
APPLICATION DATA:
APPLICATION NUMBER: US/08/234,580
CLASSIFYCATION: 435
ATTORNEY/AGENT INFORMATION: 435
NAME: MILLMAN, ROBERT A
REFERENCE/DOCKET NUMBER: 1011.1030
TELECOMMUNITY ATTORNES: 1011.1030
                                                                                                                                                                                                                                                                                                                                                                               275 AGGASATTAAAAAGCCAGTTTGAAAGCTTTGTGAAGGATATAATGTTAAACAAAGAGGAGA 334
                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 3.
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                              855 tcatgamancyatacagagatgcaacacaggagamagatccttatccttactgamatgtg 914
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                         FEATURE :
                                                                                                                           MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)371-2600
                                                                                                                                                                                                                               ACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAGCTGACCGTTAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                 CGAAGAAAGAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCTCAAATTGCGGCAC
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COMPUTER: II
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22..807
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                                                                                                                                                                                                                                                                         NAME/KEY:
COCATION:
US-08-858-197-3
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Best Local Similarity 99.6%;
Matches 683; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
975 cgaagaaagaaacagctttgaaatgcaaaaaggtgatcagaatcctcaaattgcggcac
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                                                                                       215 TCATGAAAACGATACAGAGATGCAACACAGGAGAAGATCCTTATCCTTACTGAACTGTG
                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08,
FILING DATE: APR-194
ATTORNEY/AGENT INFORMATION:
NAME: MILLMAN, ROBERT A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA
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REFERENCE/DOCKET NUMBER: 10.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -08-858-197-3
                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: METHODS FOR PROLIFERATING NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERN, KESSLER, GOLDSTEIN & FOX STREET: 100 NEW YORK AVE. N.W. SUITE 600
                                                   AGGAGATTAAAAGCCAGTTTGAÁGGCTTTGTGÁÁGGÁTÁTAATGTTAAACAAAGÁGGÁGA
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APPLICANT: CASTLE, BRIAN E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1455 caccttgcaggctgtggtggagctga 1480
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STATE: D.C.
ZIP: 20005
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22..807
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                                                                                                                                                                                              Score 681.2; DB 12
Pred. No. 8.8e-157;
0; Mismatches 3;
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RESULT 14
US-09-388-079-1
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GENERAL INFORMATION:
APPLICANT: ARUFFO, ALEJANDRO
APPLICANT: HOLLENBAUGH, DIANE
APPLICANT: LEDBETTER, JEFFREY A.
TITLE OF INVENTION: SOLUBLE LIGANDS FOR CD40
NUMBER OF SEQUENCES: 15
                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/690
FILING DATE: 31-UUL-1996
APPLICATION NUMBER: US 07/940
FILING DATE: 04-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: MISFOCK, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 5624
TELECOMMUNICATION 1270-9090
TELEPHONE: 212,790-9090
                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPBALIBLE
COMPUTER: PA PC COMPBALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
ADDITION WIMPER. IS //0/238 OFF
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TELEPHONE: 212 790-9090
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TELEX: 66141 PENNIE
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CLASSIFICATION:
                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                   STATE: New York COUNTRY: U.S.A. ZIP: 10036-2711
                                                                                                                                                                                                                                                                                                                                                                CITY: New York
STATE: New York
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RESULT 15
US-09-388-079A-1
; Sequence 1, Application US/09388079A
; GENERAL INFORMATION:
: APPLICANT: ARUFFO, ALEJANDRO
: HOLLENBAUGH, DIANI
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Best Local
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SEQUENCE CHARRCTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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99.6%;
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Pred. No. 8.8e-157;
0; Mismatches 3;
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LOCATION: 22..807
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
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Best Local Similarity
Matches 683; Conserv
                           1215 gtcaagctccatttatagccagcctctgcctaaagtcccccggtagattcgagagaatct 1274
                                                                               INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
MOLECULE TYPE: DNA (genomic)
FEATURE:
                                                                                                                                    915 aggagattaaaagccagtttgaaggctttgtgaaggatataatgttaaacaaagaggaga 974
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                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
ENTERNY APPLICATION DATA:
PPLICATION NUMBER: US/09/388,079A
PPLICATION NUMBER: US/09/388,079A
CLASSIFICATION: CONKNOWN:
NAME: MISTOCK: 31-Aug-199
ATTORNEY/AGENT INVORMATION:
REFERENCE/DOCKET NUMBER: 5624-251
TELEPHONE: 212 865-8864/9741
TELEPAX: 212 865-8864/9741
TELEPAX: 66141 DEMNITE
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: SOLUBLE LIGANDS FOR CD40 NUMBER OF SEQUENCES: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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99.6%; Pred. No. 8.8e-157;
tive 0; Mismatches 3; Indels 0;
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                                                                                                    815 CACCTTGCAGGCTGTGGTGGAGCTGA 840
                           755 GCCAAGTGAGCCATGGCATCGCTTCACGTCCTTTGGCTTACTCAAACTCTGAACAGTGT 814
                                                                 TGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAATGTGACTGATCCAA 754
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Search completed: August Job time: 5674 sec 8, 2001, 06:41:33

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Minimum DB
Maximum DB
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Perfect score:
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Listing first 45 summaries
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seq length: 2000000000
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Match Length
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1: /cgnl_7/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgnl_7/ptodata/2/pna/US06_NEW_COMB.seq:*

3: /cgnl_7/ptodata/2/pna/US07_NEW_COMB.seq:*

4: /cgnl_7/ptodata/2/pna/US08_NEW_COMB.seq:*

5: /cgnl_7/ptodata/2/pna/US09_NEW_COMB.seq:*

6: /cgnl_7/ptodata/2/pna/US00_NEW_COMB.seq:*
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US-09-808-383-5037
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US-60-278-258-10512
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Sequence 3489, Ap
Sequence 46, Appl
Sequence 123, Appl
Sequence 11351, A
Sequence 11152, A
Sequence 110472, A
Sequence 10792, A
Sequence 10792, A
Sequence 5037, Ap
Sequence 5037, Ap
Sequence 11304, A
Sequence 11304, A
Sequence 10573, A
Sequence 10573, A
Sequence 10573, A
Sequence 10773, Appl
Sequence 1077, Appl
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Sequence 1071, Appl
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855 tcatgaaa 250 tcatgaaa 250 tcatgaaat 915 aggagatt 310 aggagatt 310 aggagata 975 cgaagaaa 975 cgaagaaa 975 cgaagaaa 975 cgaagaaa 975 cgaagaaa 975 cgaagaaa 975 atgtcata 1	y Match Local Simi hes 713; 795 atggcce	28 116.4 7.5 3876 6 29 115 7.4 1429 5 30 111.4 7.2 511 5 30 111.4 7.2 516 5 31 110.6 7.1 2376 5 32 110.6 7.1 5318 6 33 110.6 7.1 2375 5 34 110.2 7.1 2375 5 35 110.2 7.1 2375 5 36 110.2 7.1 2251 6 37 109.4 7.0 5432 6 37 109.4 7.0 5432 6 39 108.8 7.0 2073 5 40 108.6 7.0 2913 6 41 108.6 7.0 2913 6 42 108.4 7.0 587 5 43 107.8 6.9 10759 6 43 107.8 6.9 10759 6 44 106.8 6.9 570 5 44 106.8 6.9 10759 6 45 105.2 6.8 3690 6 45 105.2 6.8 3690 6 45 105.2 7.0 2010 PC. GENERAL INFORMATION: NOVEL NUMBER: 0 CURRENT HYSEG, INCOMENTIAL OF INCOMENT FILING DATE: 2001-0; PRIOR APPLICATION NUMBER: 0 CURRENT FILING DATE: 2000-05-1; PRIOR FILING DATE: 2
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ascacaggagaaagatccttatc	Score 705.2; DB 1; Len Pred. No. 5.8e-163; 0; Mismatches 13; Ind gatagaagatgaaaggaatcttcatg [US-60-278-258-10748 US-09-898-888-21593 US-09-898-888-21593 US-09-849-626-1138 US-09-652-125A-9454 US-60-278-258-10899 US-09-764-864-232 US-09-808-383-5593 US-60-278-258-10596 US-09-808-383-1191 US-09-808-383-1191 US-09-808-383-91 US-09-808-383-91 US-60-278-258-10426 US-09-808-383-91 US-60-260-483-448 ALIGNMENTS ALIGNMENTS PUS0114827
ccttactgaactgtg 914	ength 1833; Idels 0; Gaps 0; Idels 0; Gaps 0; Idels 1 1 1 1 1 1 1 1 1 1	Sequence 10748, A Sequence 21593, A Sequence 1138, Ap Sequence 10761, A Sequence 10899, A Sequence 10899, A Sequence 10596, A Sequence 10596, A Sequence 10596, A Sequence 11911, App Sequence 110425, Ap Sequence 10425, A Sequence 10426, A Sequence 10, Appl Sequence 448, Appl Sequence 448, Appl Sequence 448, Appl Sequence 10426, A Sequence 10436, Appl Sequence 10426, Appl Sequence 10436, Appl Sequence 10436, Appl Sequence 10426, Appl Sequence 10436, Appl Sequence 10426, Appl Sequence

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1095 actacaccatgagcaacaacttggtaaccctggaaaatgggaaacagctgaccgttaaaa 1154

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NAME/REY: SITE
COCATION: (889)
OTHER INFORMATION: n equals
                                                                  Query Match
Best Local Similarity
Matches 657; Conserv
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LOCATION: (48)
OTHER INFORMATION: n equals a,t,g, or
LOCATION: (415)
OTHER INFORMATION: n equals a,t,g, or
LOCATION: (315)
OTHER INFORMATION: n equals a,t,g, or
NAME/KEY: STE
                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: US/09/760,481

Prior application wilder: US/09/760,481

Prior application data removed - refer to PALM or file wrapper SOFTWARE: DI NOS: 317

SEO ID NO 46

LENGTH: 950

TYPE: DNA ORGANISM: Adomo sapiens
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US-09-760-481-46
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LOCATION: (840)
OTHER INFORMATION: D
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   FEATURE.
NAME/KEY: SITE
TON: (42)
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TITLE OF INVENTION: Nucleic Acids, Proteins,
PILE REFERENCE: PTZ64
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                                                   Score 615.6; DB 5;
Pred. No. 3.8e-141;
4; Mismatches 16;
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                  SEOFTWARE: Patentin Ver. 2
SEQ ID NO 123
LENGTH: 950
TYPE: DNA
ORGANISM: HOMO Sapiens
PEATURE: SITE
LOCATION: (42)
OTHER INFORMATION: n equal NAME/KEY: SITE
LOCATION: (48)
OTHER INFORMATION: n equal NAME/KEY: SITE
LOCATION: (48)
OTHER INFORMATION: n equal NAME/KEY: SITE
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Sequence 123, Application US/09760481
GENERAL IMPORMATION:
APPLICAMY: ROSen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PTZ64
FILE REFERENCE: PTZ64
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NAME/KEY: SITE
LOCATION: (815)
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              a,t,g,
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: NAME/KEY: SITE
: LOCATION: (840)
: OTHER INFORMATION: r
: NAME/KEY: SITE
: LOCATION: (889)
: OTHER INFORMATION: r
US-09-760-481-123
Sequence 323, Application US/09760485
GENERAL INFORMATION:
APPLICANT: ROSen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PJZ17
CURRENT APPLICATION NUMBER: US/09/760,485
CURRENT FILING DATE: 2001-01-16
Prior application data removed - consult PAI
NUMBER OF SEQ ID NOS: 1477
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Pred. No. 3.8e-141;
4; Mismatches 16;
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SEQ ID NO 323
LENGTH: 950
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NAME/KEY: SITE
LOCATION: (42)
OTHER INFORMATION: |
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TYPE: DNA
ORGANISM: Homo
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                                                          gccaagtggagccatggcactggcttcacgnccttttgggtttacttcaaacttttgaac
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Pred. No. 3.8e-141;
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2822 cttggagttccag 2834
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                            2762 ggtcgagcaggcccaactggagacccaggtccttcaggtcaagcaggagaaaagggaaaa
                                                                                                                         2702
                          683 agcgggcttccag 695
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Best Local Similarity
Matches 297; Conserv
                                                                                                                                      2642 aaaggugaaaagggtgaagatggttttoccaggattcaaaggtgacatgggtotaaaaggt 2701
                                                                                                                                                                                              2582 ggatacccggggcccccgggggagtaaaggggagcagatggtgtcagagggtctcaaggggatct
                                                                                                    563 aatgetggageaggaeetgeeggaeetgeeggteeaeagggageteeaggtteeagg 622
                                                                                                                                                                                                                                                  2522 aaagaaggecagtetggagaaaagggggetetgggteeeetggteeacaaggteetatt 2581
                                                                                                                                                             503 maaggstccacaggsccccmagggagaamgaggtgsccctggtgtgscamggagscccmaggg 562
                                                                                                                                                                                                                443 ggagaggctgggcccaaaggagaagtaggtgctcctggcatgcaaggatctacaggggca 502
                                                                                                                                                                                                                                                                                                       2462 caaggaaaaocaggacttgotggacttcctggtgctgatgggcctcctggtcatcctggg 2521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using TITLE OF INVENTION: Polymoclectide Sequence Databases, and Single Nucleotide EILE REFERENCE: GX-0010-1 p-phisms Identified Thereby

CURRENT APPLICATION NUMBER: US/60/278,258

CURRENT FILING DATE: 2001-03-23

SOFTWARE: PERL PROGRAM

SOFTWARE: PERL PROGRAM

FINDMUS. 2006
                                                                                                                                                                                                                                                                                                                                                             2402 ggacctcagggtcttcctggtccacaaggtccaattggtcctcctggtgaaaaaggacca 2461
                                                                                                                                                                                                                                                                   383 aaagaaggteeetetgggaageaggggaacataggaceteaaggcaaaceaggteetaaa 442
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           2342
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ORGANISM: Homo sapiens
FEATURE:
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         143 atgitgtagcccaacagagaatggcctgcctggtcgtgatggacgggatgggagaggaggt 202
                                               gacagaggagaagttggtcaaattggcccaagaggggaagatggccctgaaggacccaaa
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3332 ccgtggcttcactggtctg 3350
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LOCATION: 50, 110, 1441, 1446
OTHER INFORMATION: a, t, C, g, or other
US-60-278-258-11152
                               3272 aggccccagaggtgacaaaggagaggctggagagcctggcgagagaggcctgaagggaca
                                                                                         3212 tgcacaaggccccatgggaccctcaggaccagctggagcccggggaatccagggtgctca 3271
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Best Local Similarity 52.4
293; Conservative
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US-60-278-258-11152
Sequence 11152, Applications
GENERAL INFORMATION:
                    691 tecagacagtgetgetetg 709
                                                                                                                                                    631 aggactcaagggggacagaggtgttcctggagacagagggaatcaaaggtgaaagcgggct 690
                                                                                                            571 agcagcaggacctgocggacctgccggtccacagggagctccaggttccagggggcccc 630
                                                                                                                                                                                                              3092 tggagtcaagggtgatcgtggtgagactggtgctgtgggagctcctggagcccctgggcc
                                                                                                                                                                       511 cacaggccccaagggagaaagaggtgcccctggtgtgcaaggagccccagggaatgctgg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL Program
LENGTH: 4840
TYPE: DNA
TYPE: DNA
                                                                                                                                                                                                                                                                                                                              2972 atotggagacagaggtoctoctggccccgtgggtcctcctggcctgacgggtcctgcagg 3031
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FEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: 410910.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Morris, MacDonald
APPLICANT: Lal, Preet1
APPLICANT: Lal, Preet1
APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polymorphisms Sequence Databases, and Single Nucleotide
FILE REFERENCE: GX-0010; Polymorphisms Identified Thereby
                                                                                                                                                                                                                                                                                                                                                                                                                   271 aggiccagitggacccaaaggagagaatggcictgciggcgaacctggaccaaagggaga 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      211 tgagaagggtgatccaggtttgccaggacctatgggggctctcagggttgcagggccctac 270
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52.48;
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Pred. No. 5.9e-23;
0; Mismatches 266;
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                                      RESULT 8
US-60-278-258-10472
Sequence 10472, Application US/60278258
GENERAL INFORMATION:
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FILE REFERENCE: Gx-0010-1 P
CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL PROGram
SEQ ID NO 10948
LENGTH: 6943
TYPE: DNA
TYPE: DNA
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Best Local Similarity
Matches 283; Conserv
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OTHER INFORMATION: Incyte ID
-60-278-258-10948
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APPLICANT: Morris, MacDonald APPLICANT: Lal, Preeti
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; OTHER INFORMATION: Incyte ID
US-60-278-258-10472
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                                                                                                                                                                                                                                      US-60-278-258-10792
Sequence 10792, Application US/60278258

GENERAL INFORMATION:
APPLICANT: Morris, MacDonald
APPLICANT: Diep, Dinh
APPLICANT: Diep, Dinh
APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Us;
TITLE OF INVENTION: Polymorphisms Identified Thereby
FILE REFERENCE: GX-0010-1 p
CURRENT APPLICATION NUMBER: US/60/278,258

CURRENT FILING DATE: 2001-03-23
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TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide
TITLE OF INVENTION: Polymorphisms Identified Thereby
FILE REFERENCE: GX-0010-1 P
CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL Program
SOFTWARE: PERL Program
SEQ ID NO 10472
LENGTH: 3753
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Best Local Similarity
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ORGANISM: Homo
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Sequence 181, Application US/09758449
GENERAL INFORMATION:
APPLICANT: Rosen et al.
ITITLE OF INVENTION: Nucleic Acids, Proteins, and
CURRENT APPLICATION NUMBER: US/09/758,449
CURRENT FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/179,065
PRIOR APPLICATION NUMBER: 60/180,628
PRIOR APPLICATION NUMBER: 60/180,628
PRIOR APPLICATION NUMBER: 10/180,628
PRIOR APPLICATION NUMBER: 10/180,628
PRIOR FILING DATE: 2000-02-04
NUMBER OF SEQ ID NOS: 1478
NUMBER OF SEQ ID NOS: 1478
SOFTMARDE: DATE: DATE: 2000-02-04
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NAME/KEY: misc_feature
OTHER INFORMATION: Incyte II
NAME/KEY: unsure
LOCATION: 7151-7153, 8062
OTHER INFORMATION: a, t, c,
US-60-278-258-10792
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SOFTWARE: Patentin Ver.
SEQ ID NO 181
LENGTH: 2060
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Best Local Similarity
Matches 284; Conserv
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SEQ ID NO 10792
LENGTH: 9033
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Pred. No. 2.8e-22;
0; Mismatches 255;
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LOCATION: (69)
OTHER INFORMATION: n e.
NAME/KEY: SITE
LOCATION: (1981)
OTHER INFORMATION: n e.
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Best Local Similarity
Matches 286; Conserv
1393 CCTCCTG 1387
                                                             1453
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                                                                                                                                                              1573 GACCGAGGGCTCCCTGGAACTCAAGGATCTCCAGGAGCAAAAGGGGGATGGGGGAATTCCT 1514
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LOCATION: (3)
OTHER INFORMATION: n equals a,t,g,
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                           agtgctg 704
                                            AAGGGTAACAAAGGCTCTACTGGACCCGCTGGCCAGAAAGGTGACAGTGGTCTTCCAGGG 1394
                                                              aagggggacagaggtgttcctggagacagagggaatcaaaggtgaaagcggggttccagac
                                                                                                     GGTCCTGCTGGTCCCTTAGGTCCACCTGGTCCTCCAGGCTTACCAGGTCCTCAAGGCCCA
                                                                                                                      99acctgccggacctgccggtccacagggagctccaggttccaggggggcccccaggactc 637
                                                                                                                                                                                                                        AAGGGACATCCTGGTTTAATTGGCCTGATTGGTCCTCCAGGAGAACAAGGGGGAAAAAAGGT
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Pred. No. 4.2e-
0; Mismatches
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Sequence 5037, Application US/09808383
GENERAL INFORMATION:
APPLICANT: Gearing, David P.
APPLICANT: Gearing, David P.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED FROM A TITLE OF INVENTION: HUMAN FETAL SKIN LIBRARY
FILE REFERENCE: 1600.1046-002
CURRENT APPLICATION NUMBER: US/09/808,383
CURRENT FILING DATE: 2001-03-13
PRIOR APPLICATION NUMBER: US 09/397,022
PRIOR APPLICATION NUMBER: US 09/397,022

RESULT 11 US-09-808-383-5037

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                   RESULT 12
PCT-US01-08656-3887/c
; Sequence 3887, Application
; GENERAL INFORMATION:
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OTHER INFORMATION: n - US-09-808-383-5037
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PRIOR FILING DATE: 1998-10-30
PRIOR APPLICATION NUMBER: US 50/107,257
PRIOR FILING DATE: 1998-11-15
PRIOR APPLICATION NUMBER: US 60/126,906
PRIOR TILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: US 60/132,099
PRIOR FILING DATE: 1999-04-30
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 5037
LENGTH: 4137
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Best Local (
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APPLICANT: Hyseq, Inc
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                                                                                                                                       2211 cctcctg 2217
                                                                                                                                                                                                                                                                                                                                                                            2031 gaccgagggctccctggaactcaaggatctccaggagcaaaagggggatgggggaattcct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         278 gttggacccaaaggagagagaggctctgctggcgaacctggacccaaagggagaaacgtgga 337
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                                                                                                                                                                                                                                         | Aagggggacagaggtttcctggagacagaggaatcaaaggtgaaagcgggcttccagac
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Pred. No. 5.3e-22;
0; Mismatches 261;
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RESULT 13
US-09-898-808-1637
US-09-898-808-1637, Application US/09898088

GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INVENTION: NOVEL CONTIGS OBTAINED FRETILE OF INVENTION: LIBRARIES
FILE REFERENCE: 20411-748CON1
CURRENT APPLICATION NUMBER: US/09/898,888
CURRENT FILING DATE: 2001-07-03
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Best Local S
Matches 281
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SOFTWARE: Custom
SEQ ID NO 3887
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: NOVEL MACROPHAGE NUCLEIC ACIDS AND POLYPEPTIDES FILE REFERENCE: 21272-066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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NAME/KEY: SIMILAR
NAME/KEY: SIMILAR
LOCATION: (3126)...(4)
LOCATION: (3126)...(4)
LOCATION: (3126)...(4)
LOCATION: 63% homologous to Homo sapiens alpha 1 (I) chain
OTHER INFORMATION: propeptide, accession number K01228, Smith-Waterman
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Local Similarity 52.5%;
hes 281; Conservative
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pred. No. 8.7e-22;
0; Mismatches 254;
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; OTHER INFORMATION: Incyte ID No: 900035.13 US-60-278-258-11304

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                                APPLICANT: MORTIS, MacDonald
APPLICANT: Lial, preet1
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polymorphisms Identified Thereby
TITLE OF INVENTION: Polymorphisms Identified Thereby
FILE REFERENCE: GX-0010-1p
CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL PROGRAM
LENGTH: 4748
TYPE: Num

LENGTH: 4748
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US-09-898-888-1637
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NAME/KEY: misc_feature
                            FEATURE:
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Best Local Similarity 52.5%;
Matches 281; Conservative
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PRIOR FILING DATE: 1999-06-28
NUMBER OF SEQ ID NOS: 45207
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 1537
LENGTH: 6191
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Pred. No. 9.6e-22;
0; Mismatches 254;
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RESULT 15
US-60-278-258-10582
Sequence 10582, Applic
GENERAL INFORMATION:
NAME/KEY: misc_feature; OTHER INFORMATION: Incyte ID No: 1099943.4; NAME/KEY: unsure; LOCATION: 3618, 3629, 3634, 3637, 4408; OTHER INFORMATION: a, t, c, g, or other US-60-278-258-10582
                                                                                                                              APPLICANT: Lal, Presti
APPLICANT: Diep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polymorphisms Identified Thereby
TITLE OF INVENTION: Polymorphisms Identified Thereby
CURRENT APPLICATION NUMBER: US/60/278,258
CURRENT APPLICATION NUMBER: US/60/278,258
NUMBER OF SEQ ID NOS: 17730
SOFTWARE: PERL Program
SEQ ID NO 10582
LENGTH: 4796
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                                                                                                                   FEATURE:
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Best Local Similarity
Matches 279; Conserv
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1526 ccccctggccccattggtaatgttggtgctcctggagccaaaggtgctcgcggcagcgct 1585
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                                                                                                                                                                                                                                  1706 ccccgtggtgagactggccctgctggacgtcctggtgaagttggtcccccttggtccccct 1765
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                                                                                                                                                                              524 ggagaagaggtgcccctggtgtgcaaggagccccagggaatgctggagcaggagcat 583
                                                                                                   584 gccggacctgccggtccacaggggctccacaggttccagggggccccccaggactcaagggg 643
                                                                                                                                                                                                                                                         464 gaagtaggtgctcctggcatgcaaggatctacagggggcaaaaggctccacagggccccaag 523
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                                                                                                                                                                                                                                                                                                                                                                                                           344 ggacctccaggacttccaggtattcctggtccagctgggaaagaaggtccctctgggaag 403
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Search completed: August 8, 2001, 05:58:16 Job time: 5722 sec

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Maximum Match 100%
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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698.8	705.2	710	710	720.2	720.2	778.6	790.4	
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89	93	97	93	93	10	94	94	. B
AF344841	HSTRAPA	HUMCD4 OL	HSCD40	HSGP39MR	AX090039	MUSSPD	BC003705	ID
AF344841 Cercocebu	X68550 H.sapiens T	L07414 Human CD40-	x67878 H.sapiens m	. 215017 H.sapiens m	AX090039 Sequence	L40156 Mus musculu	BC003705 Mus muscu	Description

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                                          NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: Robert Strausberg, Ph.D.
Tel: (301) 496-1550
Email: Robert_Strausberg@nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Sequencing Group at the Stanford Human
Center, Stanford University School of Medicine, Stanford,
                                                                                                                                                      Direct Submission
Submitted (28 FEB-2001) National Institutes of Health, Mammalian
Gene Collection (MGC), Cancer Genomics Office, National Cancer
Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
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(Dickson, Mark) mcd@paxil.stanford.edu
M., Schmutz, J., Grinwood, J., Rodrique
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Sciurognathi; Muridae; Murinae; Mus
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protein
                 Rodriquez,
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248469 B.taurus mR
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X65018 H.sapiens m
AF132496 Sus scrof
AF079105 Felis Cat
X65453 M.musculus
AF086711 Canis fam
AR044778 Sequence
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clone MGC:5668,
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TGPKGERGAPGVOGAPGNAGPAGPAGPAGPAGPAGPAGLAGPAGVPGDRGIKGES
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Pred. No. 2.6e-191;
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431 428

368 371 308 248 251 188

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 1253)
Motwani.M., White,R.A., Guo,N., Dowler,L.L., Tauber,A.I. and
Sastry,K.N.
                                                                                                                                                                                                                                                                                                   Mouse surfactant protein-D. cDNA cloning, gene localization to chromosome 14 J. Immunol. 155 (12), 5671-5677 (1995) 96094460
                                                                                                                                                                                                                                                                                                                                                                                      C-type lectin; collagenous lectin; collectin; Mus musculus (strain B6/CBAFIJ) cDNA to mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                        MUSSPD 1253 bp mRNA Mus musculus (clones 12.1, 2.1 and 4.1) (Sftp4) mRNA, complete cds.
                                                                  /gene="Sftp4"
63. .1127
                                                                                                                                                                                                   /tlssue_type="lung"
/clone_lib="lambda zap
/chromosome="14"
/gene="Sftp4"
/product="surfactant protein
                                                             NWAPGEPHNNGGAENCVETFTNGQWNDKACGEQRLVTCEF
                                                                                                                                                  /codon_start=1
                                                                                                                                                          /gene="Sftp4"
/note="SP-D"
                                                                                                                                                                                                                                      /db_xref="taxon:10090"
/clone="12.1; 4.1; 2.1"
/sex="female"
                                                                                                                                                                                                                                                                /organism="Mus musculus"
/strain="B6/CBAF1J"
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1. .1253
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                                            /gene-"Sftp4"
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                       CACCTTGCAGGCTGTGGTGGAGCTGACGCTGGGAGTCTTCATAATACAGCACAGGCTTAA
                                caccttgcaggctgtggtggagctgacgctggggagtcttcataatacagcacaggcttaa 1514
                                                                                               t999a(9agtatttgaattgcaaccaggtgcttcggtgtttgtcaatgtgactgatccaa
                                                                                      TGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAATGTGACTGATCCAA
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Mummalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 879)
Ahuja, S.U. and Bonewald, L.U.
C040 agonist compositions and methods of use
Putent: WO 0116180-A 1 08-MAR-2001;
BCARD OF REGENTS THE UNIVERSITY OF TEXAS SYSTEM (US)
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Homo sapiens
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/db_xref="taxon:9606"
193 c 190 g 22
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CGAAGAAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCTCAAATTGCGGCAC
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Hollenbaugh, D.L.
Direct Submission
Submitted (27-AUG-1992) D
Submitted (27-AUG-1992) D
                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
Submitted (21-SEP-1992) Diane L Holle
PRI Seattle, WA, 98121
The original submission [1] reported
Location/Qualifiers
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1 (bases 1 to 879)

Hollenbaugh,D., Grosmaire,L.S., Kullas,C.D., Chalupny,N.J.,
Braesch-Andersen,S., Noelle,R.J., Stamenkovic,T., Ledbetter,J.A.
and Aruffo,A.
The human T cell antigen gp39, a member of the TNF gene family, gp39 with B cell co-stimulatory activity

a ligand for the CD40 receptor: expression of a soluble form of EMBO J. 11 (12), 4313-4321 (1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRI, Seattle, WA, 9
3 (bases 1 to 879)
Hollenbaugh,D.
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                                                                                                                                                                                                                                                                                                             /product "glycoprotein 39"
/db_xref "SWISS - PROT : P29965"
/translation "MIETYNOTSPRSAATGLPISMKIFMYLLTVFLITQMIGSALFAV
/translation "MIETYNOTSPRSAATGLPISMKIFMYLLTVFLITQMIGSALFAV
YLHRRLDKIEDERNLHEDFVFMKTIORCNTGERSLSLLNCEETKSQFEGFVKDIMLNK
ETTKKENSFEMQKGODNOPQIAAHYTEEASSKTTSVLOMAEKGYYTMSNNLVTLENGKO
ETTKKROGLYYIYAOVTFCSNREASSQAPFIASLCLKSFGRFERILLRAANTHSSAKPC
GQOSIHLGGVFELOFGASVFVNVTDESQVSHGTGFTSFGLLKL"
gQOSIHLGGVFELOFGASVFVNVTDESQVSHGTGFTSFGLLKL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cell_type="PHA-activated T-cells"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
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ia; Primates;
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Pred. No. 2.3e-173;
0; Mismatches 3;
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Catarrhini; Hominidae;
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H.sapiens mRNA for C
X67878 S50586
X67878.1 GI:38411
                                                                                                                                                                                        Spriggs, M.K., Armitage, R.J., Strockbine, L., Clifford, K.N., Macduff, B.M., Sato, T.A., Maliszewski, C.R. and Fanslow, W.C. Recombinant human CD40 llgand stimulates B cell proliferation immunoglobulin E secretion
J. Exp. Med. 176 (6), 1543-1550 (1992)
                                                                                                                                                                                                                                                           Direct Submission
Submitted (28-JUL-1992) M. Spriggs,
Corporation, 51 University Street, 5
(bases 1 to 1803)
                                                                                                                                                                                                                                                                                                                   Spriggs, M.
                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutelo
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
1 (bases 1 to 1803)
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human.
                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
/cell_type="peripheral bl
46. .831
/product="CD40 ligand"
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YLHRRLDKIEDERNLHEDEVFMKTIQRCNTGERSLSLLNCEEIKSQFEGFVKDIMLNK
                                                                                                                                                              Location/Qualifiers
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Human CD40-ligand mRNA,
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L07414.1 GI:180123
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GOOSIHLGGVFELOPGASVFVNVTDPSOVSHGTGFTSFGLLKL"
112 . 183
/note="CD40 11gand"
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Pred. No. 9.6e-171;
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    ACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAGCTGACCGTTAAAA
                                                                                                                                                                                                  ATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGAT
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1 (bases 1 to 1816)
Gauchat, J.-F.M., Aubry, J.-P., Mazzei, G.J., Life, P., Jomotte, T., Elson, G. and Bonnefoy, J.-Y.
Human CD40-ligand: Molecular cloning, cellular distribution and regulation of expression by factors controlling IgE production F3BS Lett. 315, 259-266 (1992)
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/TEARSTATION-"M.ETYNOTSPRSAATGLPISMKIFMYLLTVELITOMIGSALFAV
/TLERKLDKIEDERNLHEDEVFMKTIQRCMTGERSLSLLNCEEIKSQFEGFVKDIMLNK
LEYKROSLEVIYAQVIFCSNREASSQAFFIASLCLKSPGREFILLRAANTHSSAKPC
GOOSIHLGGVFELQPGASVEVNVTDPSQVSHGTGFTSFGLLKL"
1518. 1581
/note-"dinuclectide (CA) repeat"
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/protein_id="AAA35662.1"
/db_xref="GI:180124"
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/db_xref="taxon:9606"
/cell_type="7"-lymphocyte"
/tissue_lib="lambda gt10"
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Pred. No. 9.6e-171;
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Cloning of TRAP, a ligand for CD40 on human T
Eur. J. Immunol. 22 (12), 3191-3194 (1992)
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/translation="MIETYNOTSPREAATGLPISMKIFMYLLTVFLITQMIGSALFAV
/thrridkiedernlhedfvfmktiQrcntgerslsllkoeeikSOfegfvkDimlak
EETKKENSFEMQKGDQNPOIAAHVISEASSKTTSVLOMAEKGYYTMSNNLVTLENGKQ
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Mammalia; Eutheria; Primates; C.
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1 (bases 1 to 1058)
Villinger, F., Bostik, P., Mayne.
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Submitted (02-FEB-2001) Pathology and Laboratory Medicine, Emory University School of Medicine, Winship Cancer Institute, 1365B Clifton Rd, Atlanta, CA 30322, USA
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Macaca.

E 1 (bases 1 to 1058)
(S Villinger, F., Bostik, P., Mayne, A.E., King, C.L., Genain, C.)
Weiss, W.R. and Ansari, A.A.
Cloning, sequencing and homology analysis of nonhuman prij
Fas/Fas-Ligand and co-stimulatory molecules
Immunogenetics (2001) In press
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Macaca mulatta CD154 protein mRNA, complete
AF344859.1 GI:13650015
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/product="cD154 protein"
/product="id="AAK37541.1"
/db_xref="G1:13550016"
/translation="%1550016"
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Pred. No. 6.8e-168;
Mismatches 17;
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Rat pulmonary surfactant protein (SP-D) mRNA, complete cds.
M81231
M81231. GI:207035
pulmonary surfactant-associated protein SP-D.
Rattus norvegicus (strain Fisher 344) lung cDNA to mRNA.
Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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imizu,H., Fisher,J.H.,
                                   structure of rat pulmonary surfactant protein anubi acud sequence ... Chem. 267, 1853-1857 (1992)
              Location/Qualifiers
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/organism="Rattus norvegicus"
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/product="pulmonary :
1237. .1242
/gene="SP-D"
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/db_xref="taxon:10116"
/tissue_type="jung"
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313 c 402
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/product="primary translation
/protein_id="AAA42170.1"
/db_xref="GI:207036"
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Sequence
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AR044779.
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Methods for proliferating and differentiating
density membrane CD40 ligand
Patent: US 5817516-A 3 06-OCT-1998;
Location/Qualifiers
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Pred. No. 2.1e-163;
0; Mismatches 3;
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                       ATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGAT
                                           CACCTTGCAGGCTGTGGTGGAGCTGA 840
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Aruffo,A., Hollenbaugh,D. and Ledbetter,J.A.
Soluble and its use in B cell stimulation
Patent: US 5540926-A 1 30-JUL-1996;
Location/Qualifiers
1. 840
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Pred. No. 2.1e-163;
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Best Local Similarity 99.1%;
Matches 680; Conservative
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Submitted (27-JAN-1996) A.C. Gra
Dallas, 5323 Harry Hines Blvd.,
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H.sapiens
X96710
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Grammer, A.C., Bergman, M.C., Miura, Y., Fujita, K., Davis, L.S.
                                                                                                                                                                                                                                                                        Lipsky, P.E.
The CD40 ligand expressed by human B cells
                                                                                                                                                                                                                                                    95248064 (10),
                                                                                                                                                                                                                                                                                                        EUKATYOTA; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (Dases 1 to 839)
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                                                                                                                               /organism="Homo sapiens"
/isolate="patient B.S."
/db_xref="taxon:9606"
/chromosome="xq26"
                                                                              /clone="9"
<1. .>839
                                                         /product="CD40-ligand"
181 c 180 g
                                                                                                                /cell_type="B&T"
                                                                                                 /cell_line-"HIGMXL/clone-"1"
                                                                                                                              /dev_stage="3
                                                                                                                                                                                  Location/Qualifiers
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r CD40-ligand
Score 666; DB 91;
Pred. No. 1.6e-159;
0; Mismatches 5;
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Aotus trivirgatus
Eukaryota; Metazoa; (
Mammalia; Eutheria; )
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AF344860
Direct Submission
Submitted (02-FEB-2001) Pathology and Laboratory Medicine,
University School of Medicine, Winship Cancer Institute, 1
Clifton Rd, Atlanta, GA 30322, USA
Location/Qualifiers
                                           Weiss, W.R. and Ansari, A.A.
Cloning, sequencing and homology analysis of
Fas/Fas-Ligand and co-stimulatory molecules
Immunogenetics (2001) In press
2 (bases 1 to 974)
Villinger, F.
                                                                                                  1 (bases 1 to 974)
Villinger, F., Bosti
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CD154
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Primates;
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                                                                                                   King, C.L.,
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Submitted (02-FEB-2001) Pathology and Laboratory Medicine, Emor
University School of Medicine, Winship Cancer Institute, 1365B
Clifton Rd, Atlanta, GA 30322, USA
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Villinger, F., Bostik, P., Mayne, A.E., King, C.L., Genain, C.P., Weiss, W.R. and Ansari, A.A.
Cloning, sequencing and homology analysis of nonhuman primate Pas/Fas-Ligand and co-stimulatory molecules
Immunogenetics (2001) In press
2 (bases 1 to 975)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Callitrichidae;
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Search completed: August 8, 2001, 05:54:20
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99US-0151250	-US23			Location, 22807 /*tag= z /product-		osteoblast necrosis; ectomy; lu tosis; nep sthma; lun	of	entry		DNA;		782 878	783	783	506	783	783	633	633	780	633	780	780	1878	989	964	783	929	1425	1425	786	786	786	786
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				/Qualiflers . "gp39"		cell death; inflammatory pus nephritis hritis; myosi g disease; ca	040			· BP .	ALIGNMENTS	AAQ41507 AAZ27538	AAV39001	AAV38998	AAZ27524 AAV42183	AAV12853	AAQ63960	AAZ55539	AAZ55538	AAZ55536	AAZ55545	AAZ55543	AAZ55542	AAZ55535	AAZ55540	AAV39004 AAZ55541	AAV42184	AAT58123 AAZ27537	AAZ27534	AA041516	AAV39003	AAV39002	AAV39000	AAQ63959
					apoptosis; bone loss; arthritis; esterogen loss; ; Takayasu's arteritis; tis; scleroderma;	ligand.					Murine CD40 L DNA. Mouse trimeric CD4	664	CD40 ligand gene u	Mouse CD40-L codin	CD40 ligand coding	Mouse CD40-L type	Canine mature CD15	Canine mature CD15	Canine CD154 CDNA	Feline mature CD15	Feline CD154 cDNA	Feline CD154 CDNA	Canine CD154 CDNA.	Feline CD154 cDNA.	CD40 ligand gene u	Exemplary nucleoti	CDNA encoding yeas	Human CD40-L/Fc fu	Human CD40-L/FC fu	Exemplary CD40 lig	Exemplary CD40 11g	Exemplary CD40 liq	an CD40-L type	

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence encodes a gp39 protein. It is a CD40 ligand. CD40 ligands are used for reducing osteoblast cell death or apoptosis, and for treating or preventing bone loss in animals, preferably humans, osteoporosis, or undergoing, bone loss. The bone loss is associated with cestrogen loss, esteorogensis, inflammatory arthritis, post-menopausal lupus nephritis, Takayasu's arteritis, Wegener's granulomatosis, anti-glomerular basement membrane nephritis, myositis, scleroderma, lung disease, nephrotic/nephritic syndrome, or cancer. They may also be for, an organ or bone marrow transplant.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CD40 agonist containing composition, used to reduce bone cell death or apoptosis associated with osteoporosis, osteonecrosis and inflammatory arthritis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-169007,
P-PSDB; AAB67612.
Local Similarity
                                                                    t999ag3agtatttgaattgcaaccaggtgcttcggtgtttgtcaatgtgactgatccaa
                                                                                                                           tactcagagctgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact 1334
                                                       tgggaggagtaittgaattgcaaccaggigcitcggigtitgicaaigigactgaiccaa
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Pred. No. 1.5e-188;
0; Mismatches 3;
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Sequence 1816 Bp;

520 A; 461 C;

343 G; 492 T;

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CC CD40 engagement on T cells can be used to induce T cell receptor CC engagement and enhance T cell affinity for a particular antigen. CD40 cagent, e.g. human CD40 ligand. The CD40-binding agents can also be used in methods for promoting T cell maturation, inhibiting T cell receptor altering the specificity of a T cell towards an antigen, inducing T cell reactivity towards an antigen or enhancing environmental stress-induced cell death, andigen can be inhibited, especially for a self-antigen inducing T cell cell death for antigen can be inhibited, especially for a self-antigen in an autoimmune cell disease, which includes rheumatoid arthritis, uveitis, insulin-dependent CC disease, which includes rheumatoid arthritis, uveitis, insulin-dependent CC disease mellitus, haemolytic anaemias, rheumatic fever, Crohn's disease, CC myasthenia gravis, glomerulonephritis, autoimmune hepatitis or systemic CC incus erythematosus. Inducing environmental stress-induced T cell death cCC environmental stress is a chemotherapeutic agent (claimed).
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Best Local Similarity
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               High density of differentiation; p sf9; insect cell c
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                           density membrane-bound CD40 ligand; B-lymphocyte; B-cell;
erentlation; proliferation; baculovirus; Spodoptera frugip
insect cell culture; tumour necrosis factor receptor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 1; 74pp;
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                                         atggccatagaaggttggacaagatagaagatgaaaggaatcttcatgaagattttgtat
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DB; AAR85486.
gtcaagctccatttatagccagcctctgcctaaagtcccccggtagattcgagagaatct
                                                                                    actacaccatgagcaacaacttggtaaccctggaaaatgggaaacagctgaccgttaaaa
                                                                                                actacaccatgagcaaccattggtaaccctggaaaatgggaaacagctgaccgttaaaa
                                                                                                                                atgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgggctgaaaaaggat
                                                                                                                                        atgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgggctgaaaaaggat
                                                                                                                                                                                                                               aggagattaaaagccagtttgaaggctttgtgaaggatataatgttaaacaaagaggaga
                                                                                                                                                                                                                                                                        tcatgaaaacgatacagagatgcaacacaggagaaagatccttatccttactgaactgtg
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                                                                                                                                                                                                                                                             tcatgaaaacgatacagagatgcaacacaggagaaagatccttatccttactgaactgtg
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CC The present sequence encodes human CD40 ligand which is used in the CC method of the invention. The method has been developed for proliferating CC comprises: (a) providing high density, membrane bound CD40 ligand; and CC (b) culturing one or more B cells in the presence of this ligand. The CC culture results in a proliferation in the number of B cells are CC induced to differentiate into antibody-producing cells in the presence of the B cells are CC induced to differentiate into antibody-producing cells in the presence CC of one or more cytokines. The method can be used for stimulating B-cell B-cell proliferation and activation is suppressed. Eight rounds of C division over six days can be achieved, corresponding to a 256-fold culturing methods.
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28-APR-1994;
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DB; AAW71751.
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94US-0234580.
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22..807
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/product= "CD40 ligand"
                                                                                                                                                                                                              membrane-bound
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Best Local Similarity
Matches 683; Conserv
Homo sapiens
                gp39; T-cell antigen; C:
B-cell differentiation;
                                           Genomic sequence of human gp39
                                                                   20-AUG-1994
                                                                                                     AAQ57984 standard; DNA; 840
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                CD40 ligand;
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                        B-cell
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No. 8.2e-178;
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ID AAQ5
XX AAQ5
AC AAQ5
XX 20-2
DT 20-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The complete nucleic acid sequence of human gp39 (hgp39) protein (corresp. to cDNA) and the complete AA sequence of hgp39 are presented in AAO57984 and AAR49548 respectively and conto. in plasmid CDM8-hgp39, deposited with the ATCC as E. coli, CDM8 MC1061/p3-hgp39 and assigned accession No. 69050. The human T cell antigen gp39 is a ligand for the CD40 receptor. Soluble gp39 may be produced using the expression vector CD8-gp39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Fig 1; 39pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid encoding human gp39 T cell antigen \cdot willigand for the CD40 receptor, causing proliferation and differentiation of B cells and some cancer cells
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DB; AAR59548.
                                        tactcagagctgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact
                                                                                                               gacaaggactctattatatctatgcccaagtcaccttctgttccaatcgggaagcttcga 1214
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                                               This sequence encodes a human CD40-L polypeptide which binds to CD40. CD40-L is a type II membrane polypeptide which has an extracellular region at its C-terminus, a transmembrane region and an intracellular region at its N-terminus. A soluble form of CD40-L lacks the transmembrane domain. CD40-L activity is mediated by binding with CD40 an dinduces B cell proliferation and induction of antibody secretion, including IgE. Membrane bound CD40-L acts as a CD40 agonist and soluble CD40-L acts as a CD40 antagonist. CD40-L can be used in a binding assay to detect cells expressing CD40.
                                                                                                                                                                                                                                                                                                                                                                                        New cytokine CD40-L as CD40 agonist and antagonist - is treating allergies, lupus, rheumatoid arthritis, graft-versus-host disease and insulin-dependent diabetes
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05-DEC-1991;
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region; intracellular; soluble; activity; B cell;
induction; antibody; secretion; IgE; agonist;
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Query Match Best Local Similarity

Score 657.2; DB 14; Pred. No. 3.4e-171;

Sequence

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ligand gene; I
                   /*tag- a
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DB; AAR57469.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 266 A;
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Widmer MB
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94US-0184422
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    185 C;
                                                                                                                                                                                                                                                                                                                                                                                          Score 657.2; DB 15;
Pred. No. 3.4e-171;
0; Mismatches 3;
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RESULT
AAT93782
              The present sequence represents the cDNA sequence of a novel cytokine ligand for CD40 called CD40L. CD40L is a type II membrane polypeptide, which is expressed by activated CD4+T cells, and causes B cell proliferation and induction of antibody secretion. The protein can be used to produce monoclonal antibodies, which in turn bind to CD40-expressing cells. This inhibits binding of soluble CD40 to its ligand CD40L. The monoclonal antibody against CD40L is used to inhibit proliferation of neoplastic cells, and is particularly useful in treating B-cell lymphoma (e.g. where induced after transplants or in other cases of immune deficiency such as AIDS), and also melanoma or carcinoma. Since the monoclonal antibodies inhibit neoplastic cells directly, they may not need to be coupled to a toxin or radioisotope, avoiding toxic effects on
                                                                                                                                                                                                                                                                                                                                                                                                       Treating or preventing neoplastic disease associated with CD40-expressing cells - particularly B-cell lymphoma, by administration of CD40-binding protein, preferably antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytokine; activated CD4+ T cell; CD40; neoplastic cell proliferation; B-cell
                                                                                                                                                                                                                                                                                                                                        Claim 3; Columns 19-22;
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melanoma; c
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HEALTH & HUMAN SERVICES
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Best Local Sin
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binding
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             US5961974-A
                                                                                                                 Human CD40-L coding sequence
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                                                               CD40 receptor ligand; anti-human CD40-L monoclonal antibody;
inhibitor; trimeric CD40-L; anti-Immunoglobulin M;
ral blood B cell; proliferation inhibitor; ss.
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Pred. No. 3.4
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05-DEC-1991;
23-OCT-1992;
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tgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaatgtgactgatccaa
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           tactcayagctgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact
                                         tactcagagetgeaaatacccacagttccgccaaaccttgcgggcaacaatccattcact
                                                                                   gtcaagotccatttatagccagcctctgcctaaagtcccccggtagattcgagagaatct
                                                                                                                           actacaccatgagcaacattggtaaccctggaaaatgggaaacagctgaccgttaaaa
                                                                                                                                                                       actacaccatgagcaacaacttggtaaccctggaaaatgggaaacagctgaccgttaaaa
                                                                                                                                                                                                                atgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgggctgaaaaaggat
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                                                                                                                                                                                                                                                                     aggagattaaaagccagtttgaaggctttgtgaaggatataatgttaaacaaagaggaga
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DB; AAY39938.
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91US-0805723.
92US-0969703.
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Pred. No. 3.
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This sequence covers cDNA coding for human CD40L mutein polypeptides in which the wild-type Cys residue at position 194 has been replaced by a Trp. Ser, Asp or Lys residue. The muteins, designated and c194W, C194S, C194D and C194K, are functional analogues of CD40L secretion. This property may be used to treat disorders associated muteins have a higher binding affinity for human CD40L CD40L and can be used in binding assays.

(Note: The present sequence does not appear in the specification; which is given on pages 19-20, at codon 194).
                                                                                                                                                                                                                               Armitage RJ,
                                                                                                                                                 New CD40 ligand mutein with higher CD40 affinity than native ligand - useful in binding assays, and for therapy of disorders and diseases involving low levels of B cells and antibody secretion
                                                                                                                                           Claim 1;
                                                                                                                                                                                              WPI; 1997-052320/05
P-PSDB; AAW09113, AAW09114, AAW09115,
                                                                                                                                                                                                                                                                                 07-JUN-1995;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CD40 ligand;
antibody secr
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                                                                                                                                                                                                                                         Fanslow
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retion; immunoglobulin E; cyto
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95US-0477733
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/product= CD401_mutein
/note= "Encodes CD401 in which the Cys residue
position 194 is replaced by Trp, Ser, A
or Lys"
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Best Local Similarity
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                                                                                       CD40-L
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Location/Qualifiers
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99.1%;
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No. 2.;
                                                                  melanoma;
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2.2e-170;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CD40-L is a type II membrane protein whose activity, mediated by binding of its extracellular region to CD40, includes B-cell proliferation and induction of antibody (including IgE) secretion CD40-L/CZ2 fusion protein, trimeric CD40-L (33-amino acid zipper, 8-amino acid hydrophilic sequence and the extracellular region of CD40-L), and other forms of CD40-L can be produced by recombinant methods. CD40-L cDNA has been cloned from peripheral blood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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DB; AAR76125.
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No. 7e-170;
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                                                                               A DNA fragment encoding the extracellular (soluble) region of human CD40-L was ligated to a synthetic oligonucleotide sequence coding for a leader peptide, a 33 amino acid leucine zipper sequence (AAR53968) and the Flag (RTM) linker sequence. Cells expressing the fusion construct are grown to accumulate oligoneric, soluble CD40-L trimerises in solution and fusion proteins comprising the trimerises in solution and fusion proteins comprising
                                                  Sequence
                                                                                                                                                                                             Prepn. of soluble oligomeric mammalian proteins - using cells to express a fusion protein comprising a leucine a domain and a heterologous mammalian protein
                                                                                                                                                                            Example 1; Page 22-23; 35pp; English.
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13-AUG-1993;
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93US-0107353.
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/product= human_CD40-L
/note= "nucleotides 148-783 code for the
extracellular region (amino acids 50-261)"
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Pred. No. 9.8e-169;
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                      01-DEC-1997;
09-DEC-1996;
(REGC ) UNIV CALIFORNIA
                                                     08-DEC-1997;
                                                                                                                           CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; autoimmune disorder; rheumatoid arthritis; vaccine; ss.
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                                                                         18-JUN-1998
                                                                                            W09826061-A2
                                                                                                                                                                    CD40 ligand gene used in the course of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
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Chimeric - Homo sapiens.
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The present sequence represents an exemplary CD40 ligand gene, comprising nucleotides encoding the extracellular domains (Domains III and IV) and transmembrane domain (Domain II) of human CD40 ligand gene
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                                                                                                                             Disclosure;
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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1: /cgnl_7/ptodata/1/ina/5A_COMB.seq:*
2: /cgnl_7/ptodata/1/ina/5B_COMB.seq:*
3: /cgnl_7/ptodata/1/ina/6A_COMB.seq:*
4: /cgnl_7/ptodata/1/ina/6B_COMB.seq:*
5: /cgnl_7/ptodata/1/ina/backfilesl.seq:*
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US-08-431-055-3
US-08-144-422-7
US-08-249-189-11
US-08-484-624A-11
US-08-484-624A-12
US-08-484-624A-13
US-08-484-624A-20
US-08-484-624A-11
US-08-484-624A-20
US-08-484-624A-20
US-08-484-624A-20
US-08-484-624A-20
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US-08-484-624A-20
US-08-484-624A-20
US-08-484-624A-1
US-08-484-624A-1
US-08-484-624A-2
US-08-484-624A-2
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Query Match 43. Best Local Similarity 99. Matches 683; Conservative

43.98;

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Score 681.2; DB 1; Pred. No. 8.5e-178; 0; Mismatches 3;

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Gaps

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    STREET: 100 NEW YORK AVE. N.W. SUI-
CITY: WASHINGTON
STATE: D.C.
ZIP: 20005
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
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Patent No. 5817516
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8-431-055-3
                                                                            NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERN, KESSLER, GOLD
STREET: 100 NEW YORK AVE. N.W.
                                                                                                       **NERAL INFOFMATION: MERİLYN R
APPLICANT: KEHRY, MERİLYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING
TITLE OF INVENTION: DIFFERENTIATING B CELLS, A
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   APPLICATION NUMBER:
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             Release #1.0, Version
US/08/431,055
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COCATION:
US-08-431-055-3
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Best Local Sin
Matches 683;
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Matches
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/234
FILING DATE: 28-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: MILLMAN, ROBERT A
REGISTRATION NUMBER: 36,217
REFERENCE/DOCKET NUMBER: 1011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)371-2540
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDENNESS: hoth
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FEATURE:
STRANDEDNESS: bot
                                                                         TGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAATGTGACTGATCCAA
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                                               tgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaatgtgactgatccaa 1394
                                                                                                                                                              GACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAATCGGGAAGCTTCGA
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Pred. No. 8.5e-178;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/690,096
FILING DATE: 31-TUL-1996
CLASSIFICATION: 536
PRIOR APPLICATION: DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 212 790-9090
TELEFAX: 212 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 562
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
TELEFAX: 212 600-0000
                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/940,605
FILING DATE: 04-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                 915
                                                                                                              855
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LOCATION:
                                                                                                                                                                                                                                                    Local Similarity
nes 683; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
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CGBABGABABACAGCTTTGABATGCABABAGGTGATCAGBATCCTCABATTGCGGCAC
                                                                  aggagattaaaagccagtttgaaggctttgtgaaggatataatgttaaaccaaagaggaga 974
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                                                AGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTAAACAAAGAGGAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HOLLENBAUGH, DIANE
LEDBETTER, JEFFREY A.
VENTION: SOLUBLE LIGANDS FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ARUFFO, ALEJANDRO
                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pennie & Edmonds
                                                                                                                                                                                                                                                                                                                                                                                                     DNA (genomic)
                                                                                                                                                                                                                                                                 43.98;
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                                                                                                                                                                                                                                                Score 681.2; DB 2;
Pred. No. 8.5e-178;
D; Mismatches 3;
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                                                                                                                                                                                                                                                                               Length 840;
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RESULT 4
US-08-184-422-7
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                               APPLICATION NUMBER: 08/009,25
FILING DATE: 01/22/93
ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PATRICIA ANNE
REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                            SOFTWARE: MS Word for Apple 5.1, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: FANSLOW, WILLIAM
APPLICANT: RENSHAW, BLAIR
APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDMER, MICHAEL
TITLE OF INVENTION: DETECTION AND TREATMENT OF
TITLE OF INVENTION: IN A CD40 LIGAND GENE
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: Apple OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPERATING SYSTEM: APPLE OPE
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REFERENCE/DOCKET NUMBER: 28
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                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tactcagagctgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact 1334
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                               2810-A
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 840 base pairs

7:

TELEFAX: TELEPHONE:

2065870430 2065870606

LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single

US-08-184-422-7

NAME/KEY: CDS LOCATION: 46..831

ORIGINAL SOURCE:
ORGANISM: Homo sapiens
IMMEDIATE: SOURCE:
CLONE: CD40-L
FEATURE:

MOLECULE TYPE: CI HYPOTHET:CAL: NO ANTI-SENSE: NO

TOPOLOGY:

linear CDNA

ry Match 42.3%; It Local Similarity 99.5%; ches 659; Conservative

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915 aggaçattaaaagccagtttgaaggctttgtgaaggatataatgttaaacaaagaggaga 974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          299 AGGACATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTAAACAAAGAGGAGA
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                                                           tgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaatgtgactgatccaa 1394
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Pred. No. 3.3e-171;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                               ; NAME/KEY:
; LOCATION:
US-08-360-923A-1
                                                                                                                                                                                                                                                                                                                                      NAME: PETKINS, PATRICIA A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2818
TELECOMMUNICATION INFORMATION:
TELEPHONE: (2065)587-0430
TELEPAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                            Query Match
Best Local Similarity
Matches 659; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: Macintosh
OPERATING SYSTEM: Apple Macintosh System 7.:
SOFTWARE: Microsoft Word for Macintosh, Ver.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/360,923A
FILING DATE: December 21, 1994
CLASSIFICATION: 424
PRIOR APPLICATION 127A:
APPLICATION NUMBER: USSN 08/172,664
FILING DATE: December 23, 1993
TARSTETCATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-360-923A-1
                                                                                                                                                                                                                                                                               MOLECULE TYPE: CDI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1,
Patent No. :
                                                                                                                                                                                                                      CLONE: CD40-L
FEATURE:
239 TCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTG
                                                ORIGINAL SOURCE:
ORGANISM: Hom
                                                                                                                                                                                                                                            IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: LONGO, DAN L.
APPLICANT: MURPHY, WILLIAM
TITLE OF INVENTION: METHOD OF PREVENTING OR TREATING
TITLE OF INVENTION: EXPRESSING CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 51 UNIVERSITY: SEATTLE STATE: WASHINGTON
             tcatgaaaacgatacagagatgcaacacaggagaaagatccttatccttactgaactgtg 914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     , Application US/08360923A
5674492
                                                                                                              Conservative
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                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                      linear
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FANSLOW, WILLIAM
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                                                                                                                                                                                                                                                                                                                                  single
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                                                                                                                       42.3%;
99.5%;
                                                                                                       Score 657.2; DB 1;
Pred. No. 3.3e-171;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                    2818-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BY NEOPLASTIC CELLS
                                                                                                                              Length 840;
                                                                                                       Indels
                                                                                                      0;
                                                                                                 Gaps
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1035

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В Ş 밁 ρy 밁 δÃ В Ş В δÕ

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US-08-249-189-11
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: Apple MacIntosh
COMPUTER: Apple MacIntosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft word for Apple, version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/249.189
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION MADER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION: 514
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                 APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANLE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                                                                                                   ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
CITY: SEATTLE
STATE: WASHINGTON
                                                                                                                                                                                                                     COUNTRY: U
ZIP: 98101
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Best Local Similarity 99.5%;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENCTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLONE:
FEATURE:
              1275
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ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PATTICIA A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
FILING DATE: October
                                                                                                                               1155
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HYPOTHETICAL: NO
ANTI-SENSE: NO
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CLONE: CD40-L
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LOCATION: 46..831
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FILING DATE: December
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
tactcagagctgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact 1334
                                       gacaaggactctattatatctatgcccaagtcaccttctgttccaatcgggaagcttcga
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N: 514
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ber 5, 1991
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APPLICATION NUMBER: US/08/484,624A
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,733
FILING DATE: June 07, 1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION NUMBER: 07/783,707
FILING DATE: December 5, 1991
CLASSIFICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
CLASSIFICATION:
APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
CLASSIFICATION:
APPLICATION SATA:
APPLICATION NUMBER: 34,693
REFERENCE, TORCKET NUMBER: 34,693
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REFERENCE, TORCKET N
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US-08-484-624A-11
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INFORMATION FOR SEQ ID NO:
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                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: E40 base pairs
TYPE: nucleic acid
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APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MOGREW, JEFFERY
TITLE OF ENVENTION: NOVEL CYTOKINE THAT BI
CORRESPONIZENCE ADDRESS:
ADDRESSIES.
                                                                MOLECULE TYPE:
                                                                                   TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: line
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, ve
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CITY: S
STATE:
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                                                                                      linear
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                                                                                                         single
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Sequence 11, Application U
Patent No. 5981724
GENERAL INFORMATION:
APPLICANT: ARMITAGE, R
APPLICANT: FANSLOW, WI
APPLICANT: SPRIGGS, ME
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US-08-477-733B-11
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Best Local Similarity 99.5%;
Matches 659; Conservative
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                                                                                                                               479
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                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                               ORGANISM: Homo
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                           NAME/KEY:
LOCATION:
CA 840
                                                                                                                                                                                                                                                                                                          CLONE: CD40-L
                                                          TACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAACAATCCATTCACT
                                                              tactcagagctgcaaatacocacagttccgccaaaccttgcgggcaacaatccattcact 1334
                                                                                                   CGAAGAAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAATCCTCAAATTGCGGCAC
CGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAATCCTCAAATTGCGGCAC
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Pred. No. 3.
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ARMITAGE, RICHARD FANSLOW, WILLIAM SPRIGGS, MELANIE

US/08477733B

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; NAME/KEY: CDS
; LOCATION: 46.
US-08-477-733B-11
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                                                                                                               Query Match
Best Local Similarity
Matches 659; Conserv
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/969,703
FILING DATE: OCCODER 23, 1992
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 2065870430
TELEFAX: 2065870606
NFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,733B
FILING DATE: June 07, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802-D
                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                          ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: I
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APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: NOVEL CYTOKINE NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                       CLONE: CD40-L
855 tcatgamaacgatacagagatgcmacacaggagamagatccttmtccttmctgmactgtg
                                           179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/783,707 FILING DATE: October 25, 1991 CLASSIFICATION: 435
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                                                            atggccatagaaggttggacaagatagaagatgaaaggaatcttcatgaagattttgtat 854
                                       ATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCATGAAGATTTTGTAT 238
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51 UNIVERSITY STREET
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MORRIS, ARVIA E.
MCGREW, JEFFERY
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                                                                                                                   Conservative
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                                                                                                           Score 657.2; DB 2;
Pred. No. 3.3e-171;
Pred. No. 3.3e-171;
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                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/08763995
Patent No. 6017527
GENERAL INFORMATION:
                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Power Macintosh 7200/90
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Macintosh, Versior
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                 APPLICANT: MARASKOVSKY, EUGENE
TITLE OF INVENTION: METHOD OF A
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN
FILING DATE: 10 JUL 1996
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                                                              FILING DATE:
CLASSIFICATION:
                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                         ADDRESSEE: IMMUNEX CORPORATION STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tactcagagctgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact 1334
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                                                                                                                                                                                                                                                                                                                      INFORMATION:
                                                                                                                                                                                      98101
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INFORMATION:
                                                                                          US/08/763,995
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                                                                                                                     Version #6.0.1
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REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2845.
TELECOMMUNICATION INFORMATION:
TELEPAX: (206)587-0430
TELEPAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 42.3
Best Local Similarity 99.3
Matches 659; Conservative
        1395 gccaag:gagccatggcactggcttcacgtcctttggcttactcaaactctgaacagtgt 1454
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ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                    419 ATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGAT
                                                                                                                                                                                                                                                                                                          359
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CLONE: CD40-L
FEATURE:
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                                      tactcagagetgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact 1334
GCCAAGTGAGCCATGGCACTGGCTTCACGTCTTTGGCTTACTCAAACTCTGAACAGTGT
                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                            CGRAGAAGAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCTCAAATTGCGGCAC 418
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838
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TELEPHONE: 2065870430
TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
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US-09-088-913A-11
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                                                                                                   TOPOLOGY: linear MOLECULE TYPE: CDN HYPOTHETICAL: NO ANTI-SENSE: NO
                                    CLONE: CD40-L
FEATURE:
                                                             IMMEDIATE SOURCE:
                                                                                           ORIGINAL SOURCE:
                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 34,693
TELECOMMUNICATION INFORMATION;
TELEPHONE: 205870430
TELEPAX: 205870430
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, version
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/477,733
FILING DATE: June 07, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: 07/969,703
NAME/KEY: CDS
LOCATION: 46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/088,913A FILING DATE: CLASSIFICATION: DATA: PRIOR APPLICATION DATA:
                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: GIBSON, MARYLOU
APPLICANT: MORRIS, ARVIA E
APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSE: IMMI
STREET: 51 UNIV
CITY: SEATTLE
STATE: WASHINGT
COUNTRY: USA
ZIP: 98101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                               840 base pairs
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USA
                                                                             Homo sapiens
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51 UNIVERSITY STREET
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GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: DAVISON, BARRY
APPLICANT: FANSLOW, WILLIAM
APPLICANT: FENSHAW, BLAIR
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: WIDMER, MICHAEIE
APPLICANT: TANGUNG
UNITED OF INVENTION: DEFECTIVE CD40L (as amended)
UNMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESSES:
ADDRESSES: TAMGUNG CORRESPONDENTION
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    STREET:
                            ADDRESSEE:
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SEATTLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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CURRENT APPLICATION DATA:

APPLICATION UNMBER: US/08/589,771B
FILING DATE: January 22, 1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/009,258
FILING DATE: 01/22/93
CLASSIFICATION UNMBER: 08/009,258
FILING DATE: 01/22/93
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: HENRY, JANIS C.
REGISTRATION NUMBER: 34,347
REFERENCE/DOCKET NUMBER: 2810-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPAX: 2065870430
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US-08-589-771B-7
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Best Local Similarity
Matches 659; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: MS-DOS/Windows 95
SOFTWARE: Word for Windows 95, 7.0a
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IMMEDIATE SOURCE:
CLONE: CD40-7
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ORGANISM: Hom
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HYPOTHETICAL: NO
ANTI-SENSE: NO
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atgtcataagtgaggccagcagtaaaacaacatctgtgtttacagtgggctgaaaaaggat 1094
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                                                                                      aggagattaaaagccagtttgaaggctttgtgaaggatataaatgttaaaccaaagaggaga
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llarity 99.5%;
Conservative
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TELEPHONE: (206)587-0430
TELEPHONE: (206)233-0644
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 786 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                    COMPUTER EADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, Version 5.
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/446,922
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: USSN 08/107,353
FILING DATE: 08-13-93
CLASSIFICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION 435
PRIOR APPLICATION STATE: 08-13-93
CLASSIFICATION: 435
PRIOR APPLICATION: TATE APPLICATION: 435
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GENERAL INFORMATION:
                                                                                           ORIGINAL SOURCE:
                                                                                                                    HYPOTHETICAL: 1
                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSIE: Immunex Corporation
STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: CDS
                                           STRAIN: CD40-L
                                                                         ORGANISM:
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                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                STRANDEDNESS:
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RESULT 13
PCT-US93-10034-3
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 650; Conserv
                                                                                                 APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                 STREET: 51 Uni-
CITY: Seattle
STATE: WA
                                                                                                                                                                                                                                      1335
                                                                                                                                                                                                                                                                                                                                                             1275
                                                                                                                                                                                                                                                                                                                                                                                 1215 gtcaagctocatttatagccagcctctgcctaaagtcccccggtagattcgagagaatct 1274
                                                                                                                                                                                                                                                                                              674
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                                                COUNTRY: U
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                                                                            ADDRESSEE: Immunea C. STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                            494
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          795 atggccatagaaggttggacaagatagaagatgaaaggaatcttcatgaagattttgtat
                                                                                                                                                                                                                                                                               GTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGATTCGAGAGAATCT
                                                                                                                                                                                                                                                                                                                                                                                                               cgaagaaagaaaacagctttgaaatgcaaaaaggtgatcagaatcctcaaaattgcggcac 1034
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                                                         USA
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99.5%; Pred. No. 9.5
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9.5e-169;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (206)587-04.
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                           1095
                                                                                                                                   1215
                                                                                                                                                                                                                                                                                               1035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 786 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 1003
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE:
                                                                                                                                                                                                                434
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                                                                                                                                                              494
                                                                                                                                                                                                                                                                    374
                                                                                                                                                                                                                                                                                                                   194 TCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: CDS
LOCATION: 1..783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Human
STRAIN: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: si
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                     gtcaagctccatttatagccagcctctgcctaaagtcccccggtagattcgagagaatct 1274
                                                                                                                                                                        gacaaggactctattatatctatgcccaagtcaccttctgttccaatcgggaagcttcga 1214
                                                                                                                                                                                                               ACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAGCTGACCGTTAAAA 493
                                                                                                                                                                                                                            tgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaatgtgactgatccaa 1394
                                                   tactcagagctgcaaatacccacagttccgccaaaccttgcgggcaacaatccattcact 1334
                                                                                                                                                            GACAAGGACTCTATTATATCTATGCCCCAAGTCACCTTCTGTTCCAATCGGGAAGCTTCGA
                                                                                                                                                                                                                                                                                                                                                                                                                                          tcatgaaaacgatacagagatgcaacacaggagaaagatccttatccttactgaactgtg 914
                                                                                                         GTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGATTCGAGAGAATCT
                                                                                                                                                                                                                                                                   ATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGAT 433
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Pred. No. 9.5e-169;
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RESULT 14
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                                                                                                                                                                                                                                                                                                     TELEPHONE: 2065870430
TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A,
REGISTRATION NUMBER: 34.693
REFERENCE/DOCKET NUMBER: 2807
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/249,189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 1425 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: SRINIVASAN, SUBHASHINI APPLICANT: GIBSON, MARYLOU TITLE OF INVENTION: NOVEL CYTOKINE
                                                                                                                                                                          HYPOTHETICAL: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 07/80 FILING DATE: December 5, CLASSIFICATION: 514 PRIOR APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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                                 FEATURE:
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NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION: 514
                                                                                                                                             CLONE:
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ER: 2802-C
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US-08-249-189-15

Query Match
Best Local Similarity
Matches 647; Conserv

Conservative

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Indels Length

Gaps

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1425; 0;

41.18; 97.98;

Score 638.6; DB 2; Pred. No. 5.5e-166; Mismatches

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-787 gtt::cctgatggccatagaaggttggacaagatagaagatgaaaggaatcttcatgaaga 846

GTCCGGAGGCGGGGGTTCAAGTTCTGACAAGATAGAAGATGAAAGGAATCTTCATGAAGA 824

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RESULT 15
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                          APPLICANT: ARWITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS
CORRESPONDENCE ASSESSED
          CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX C
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NAME/KEY: Sig_peptide

LOCATION: 4..78

US-08-484-624A-15
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Best Local Similarity
Matches 647; Conserv
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INFORMATION FOR SEQ ID NO: 15
SEQUENCE CHARACTERISTICS:
LENGTH: 1425 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                                                        NAME/KEY:
LOCATION:
FEATURE:
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907 gaactgtgaggagattaaaagccagtttgaaggctttgtgaaggatataatgttaaacaa
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/
FILING DATE: June 07, 1
CLASSIFICATION:
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PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/783,707
FILING DATE: October 25, 1991
CLASSIFICATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: Decembe
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LOCATION:
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APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,624A
                                        Conservative
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Pred. No. 5.5e
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Search completed: August 8, 2001, 05:07:49 Job time: 7993 sec

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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ACCESSION
VERSION
KEYWORDS
SOURCE
ORGANISM
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BG518956
LOCUS
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TITLE
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Mammalia; Eutheria; Rodentia; Sciurognath
1 (bases 1 to 846)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                      BG518956.1
EST.
                                                                 mRNA sequence.
BG518956
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                                                house mouse.
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                              Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                            musculus
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            Gene Collection (MGC)
                                                                            clone IMAGE: 3481958 5',
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AI326642 mo02a01.y
AA096905 mo02a01.r
BF599437 263218 MA
BG542824 602571588
BG53762 602565729
BB606038 BB606038
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BG072353 H3109E10-
AW354680 36752 MAR
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BF119140 601755433
AK019448 Mus muscu
BE290460 601088534
BF535428 602051822
BE307855 601096339
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BE308199 601091376
BE916416 6010677002
BG482911 602502918
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Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humphre
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM8510 row: o column: 15
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/clone="IMAGE:3481958"
/clone=lib="NCI_CGAP_Mam5"
/tissue_type="tumor, gross t
/dev_stage="7 months"
/lab_host="DH10B"
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/strain="C57/B6"
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National Institutes of Health, Mammalian Gene Collection (MGC)
L Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by:Incyte Genomics, Inc.
Clone distribution: NCI-CGAP Clone distribution information of found through the I.M.A.G.E. Consortium/LLNL at:
Plate: LLAM9260 row: p column: 07
High quality sequence stop: 745.
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Eutheria; Rodentia;
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Site_2: SalI; transgenic model WNT-1, expression driv
MMTV-LTR enhancer; Cloned unidirectionally. Primer:
dr. Library constructed by Life Technologies.
Investigator providing samples: Gilbert Smith, NIH*
                                                                                                                                                                                                                                                 /organism="Mus musculus"
/strain="CZECH II"
/db_xref="taxon:10090"
/clone="IMAGE:4015350"
/clone_lib="NCI_CGAP_Lu30"
/tissue_type="tumor, metastatic to mammary"
/lab_host="DH10B"
                                                                                                                 40.78;
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                                                                                               Score 632; DB 145;
Pred. No. 3.4e-166;
0; Mismatches 15;
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Sciurognathi; Muridae; Murinae; Mus.
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L Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humphr.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
http://image.llnl.gov
Plate: LLAM9311 row: m column: 02
High quality sequence stop: 673.
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                   GAGCCCCAGGGAATGCTGGAGCAGCAGGAC---TGCGGGAACTGCGGTCCACAGGAAGCTC
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/clone_lib="NCI_CGAP_Mam5"
/tissue_type="tumor, gross tissue"
/dev_stage="7 months"
/lab_host="DH10B"
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Pred. No. 1.9e-163;
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                          ccctgggaaatctgggagcagaaatgaagagcctctcgcagagatcagtacccaacacct 131
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EST.
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Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM10334 row: h column: 21
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National Institutes of Health, Mammalian
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Site_2: NotI; Cloned unidirectionally. Primer: Oligo-
Library constructed by Life Technologies. Investigator
providing samples: Gilbert Smith, NIH*
202 c 266 g 142 t
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Pred. No. 3e-162;
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                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM9177 row: b column: 15
High quality sequence stop: 683.
Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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                   1.90
/note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: Saiste_2: NotI; Cloned unidirectionally. Primer: Oligo clibrary constructed by Life Technologies. Investigator providing samples: Gibert Smith, NIH"

174 C 254 g 121 t
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/dev_stage="3 months, virgin"
/lab_host="DHIOB"
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Tissue Procurement: Gilbert Smith,
cDNA Library Preparation: Life Tec
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                       NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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                                                        Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; 1 (bases 1 to 639)
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mRNA sequence.
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Pred. No. 2.7e-158;
0; Mismatches 7;
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Sciurognathi; Muridae; Murinae; Mus
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Best Local Similarity 99.2%;
Matches 600; Conservative
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                                                         Bagbaggtccctctgggabgcaggggabcbtaggacctcaaggcabaccaggtcctaaag
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CCCC
                                                                                                                      AAGGCTCCACAGGCCCCAAGGGAGAAAGAGGTGCCCCTGGTGTGCAAGGAGCCCCAGGGA
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Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM8787 row: i column: 13
High quality sequence stop: 589.
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603
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/db_xref="taxon:1090"
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/clone_!ib="NCI_CGAP_Mam1"
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/dev_stage="3 months, virgin"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note**Organ: mammary; Vector: pCMV-SPORT6; Site_1: Sa Site_2: NotI; Cloned unidirectionally. Primer: Oligo Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
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Pred. No. 2.2e-153;
D; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 166;
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                                                                                                                                                                                                                                                      AATGAAGAGCCTCTCGCAGAGATCAGTACCCAACACCTGCACCCTAGTCATGTGTAGCCC
                                                                                                                                                                                                                                                                        aatgaagagcctctcgcagagatcagtacccaacacctgcaccctagtcatgtgtagccc 153
tggactaagtggacctccaggacttccaggtattcctggtccagctgggaaagaaggtcc
                                                  tccagttggacccaaaggagagaatggctctgctggcgaacctggaccaaagggagaacg
                                                                                                                                   gaagggtgatccaggtttgccaggacctatggggctctcaggggttgcagggccctacagg
                                                                                                                      GAAGGGTGATCCAGGTTTGCCAGGACCTATGGGGGCTCTCAGGGTTGCAGGGCCCTACAGG
                                                                                                                                                                                      AACAGAGAATGGCCTGCCTGGTCGTGATGGACGGGATGGGAGAGAAGGTCCACGGGGTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-remail.nih.gov
Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humphre
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MCC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://lnage.llnl.gov
Plate: LLAM8510 row: o column: 15
High quality sequence stop: 569.
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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1 (bases 1 to 571)
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                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
/strain="C57/B6"
/db_xref="taxon:10090"
/clone="IMAGE:3481958"
/clone_1ib="NCI_CGAP_Mam5"
/tissue_type="tumor, gross t.
/dev_stage="7 months"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="Organ: mammary; Vector: pCMV-SPORT6; Site_1: Sal:
Site_2: NotI; Cloned unidirectionally. Primer: Oligo do
Library constructed by Life Technologies. Investigators
providing samples: Lothar Hennighausen/Robin Humphreys,
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Rodentia;
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                                                                                                                                                                                                                                                                                                                                                                                                                        166; Length 571;
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Best Local Similarity 99.4
Matches 534; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MCC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM8757 row: 1 column: 19
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National Institutes of Health, Mammalian
Unpublished (1999)
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Manmalia; Eutheria;
1 (bases 1 to 537)
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BE370239
BE370239.1 GI:9315602
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     house mouse.
                                                                                                                                                                                            137
                                                                                                                                                                             /note="organ: lung; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo;
Library constructed by Life Technologies. Investigator
providing samples: Gilbert Smith, NIH*
a 130 c 182 g 88 t
                                                                                                                                                                                                                                                          /Organism="Mus musculus"
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/GLONE="TAXON:10900"
/Clone="IMAGE:3590106"
/Clone=lib="NCI_CGAP_Lu29"
/tissue_type="spontaneous tumor, metastatic to Stem cell Origin."
                                                                                                                                                                                                                                                                                                                                                                                           ocation/Qualifiers
                                                                                                               34.3%;
                                                                                                 0;
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                                                                                          Score 532.2; DB 166;
Pred. No. 3e-138;
0; Mismatches. 3; I
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Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                cDNA Library Preparation: Life Technologies, Inc. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC Clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: Plate: LLAMI0329 row: d column: 10
                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith,
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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Eukaryota; Metazoa;
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3812
/note="Organ: mammary; Vector: pCMV-SPORT6;
Site_2: NotI; Cloned unidirectionally. Prim
Library constructed by Life Technologies. In
providing samples: Gilbert Smith, NIH*
                                                   /clone="IMAGE:4486233"
/clone=lib="MCI_CGAP_Mam1"
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/dev_stage="3 months, virgin"
/lab_host="DH10B"
                                                                                                         /organism="Mus musculus"
/strain="FVB/N"
/db_xref="taxon:10090"
                                                                                                                                                            ocation/Qualifiers
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Rodentia;
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Sciurognathi; Muridae; Murinae; Mus
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CDNA clone IMAGE:4486233 5'
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae;
1 (bases 1 to 662)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collect
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov plate: LLAMB767 row: o column: 03 High quality sequence stop: 614. Location/Qualifiers
                                                            Email: cgapbs-remail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MCC clone distribution information can
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162 c 230 g 107 t
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/clone_lib="NCI_CGAP_Mam1"
/tissue_type="tumor, biopsy sample"
/dev_stage="3 months, virgin"
/lab_bost="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /strain="FVB/N"
/db_xref="taxon:10090"
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GGTCCAGCTGGGAAAGAAGGTCCCTCTGGGAAGCAGGGGAACATAGGA-CTCAAGGCAAA 425
                                                                              gaacctggaccaaagggagaacgtggactaagtggacctccaggacttccaggtattcct 370
                                                              GAACCTEGACCAAAGGGAGAACGTGGACTAAGTGGACCTCCAGGACTTCCAGGTATACCT
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics; Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Phttp://image.llnl.gov
Plate: LLAM10256 row: n column: 22
H1gh quality sequence stop: 630.
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Site_2: NotI; Cloned unidirectionally. Primer: Oligo c
Library constructed by Life Technologies. Investigator
providing samples: Gilbert Smith, NIH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="IMAGE:4458453"
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0; Mismatches 72; Indels 14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cyapbs-remail.nih.gov

Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humphre;
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
CNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can ]
found through the I.M.A.G.E. Consortium/LLNL at:
http://mage.llnl.gov
Plate: LLANS 512 row: Column: 08
High quality sequence stop: 547.
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NHH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 653)
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                                                                                              /organism="Mus musculus"
/strain="57/B6"
/db_xref="taxon:10000"
/clone="IMAGE:3482431"
/clone_1ib="NCI_CGAP_Mam5"
/tissue_type="tumor, gross tissue"
/dev_stage="7 months"
/lab_host="DH10B"
                        providing NIH*
                         /note-"Organ: mammary; Vector: pCMV-SPORT6; Site_1: Sall Site_2: NotI; Cloned unidirectionally. Primer: Oligo d7 Library constructed by Life Technologies. Investigators providing samples: Lothar Hennighausen/Robin Humphreys,
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                     Email: cgapbs-remail.nih.gov
Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humphr
Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humphr
CDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information car
found through the I.M.A.G.E. Consortium/LLNL at:
http://lmage.llnl.gov
Plate: LLAM8521 row: h column: 20
High quality sequence stop: 554.
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collectionships (1000)
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BE308199
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601091376F1 NCI_CGAP_Mam5
                                                                                                                                                          Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                       Mus musculus
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          quality sequence stop: 554.
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M Mus musculus
Eukaryota; Metazoa; Chordata; Cranlata; Ver
Mammalla; Eutherla; Rodentla; Sclurognathi;
1 (bases 1 to 835)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Ge
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
                                                                                                                                                                                                                                                                                            tcagaaagctgcattgttccctgatggccatagaaggttggacaagat 819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    aggtgcccctggtgtgcaaggagccccagggaatgctggagcagcaggacctgccggacc
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                                                                                                                                                                                                                                                                                                                                                     GCAGCAGATGGAGGCCTTAAAAGGAAAACTACAGCGTCTAGAGGTTGCCTTCTCCCACTA
                                                                                                                                                                                                                                                                                                                                                                                                            TGTTCCTGGAGACAGAGGAATCAAAGGTGAAAGCGGGCTTCCAGACAGTGCTGCTCTGAG
                                                                                                                                                                                                                                                                                                                                                                                                                          tgttcctggagacagaggaatcaaaggtgaaagcggggcttccagacagtgctgctctgag
                                                                                                                                                     mRNA sequence.
BE916416
BE916416.1 GI:
EST.
                                                                                                                                                                                                           BE916416 835 bp mRNA
601667002F1 NCI_CGAP_Mam1 Mu
                                                                                                                                           house mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculu:
/strain="C57/B6"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="NCI_CGAP_Mam5"
/tissue_type="tumor, gross
/dev_stage="7 months"
/lab_host="DH10B"
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Pred. No. 3.6e-116;
D; Mismatches 9; I
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                                                                                                Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                               musculus
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IMAGE:3967032
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Murinae; Mus
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Best Local
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CCA-GGAATGCTGGAGCAGCAGGACCTGGCGGACCTGCCAGGTCCACAGGGAGCTCCAGGT
                                                                                                           GGGCAAAAGGCTCCACAGGCCCCAAGGGAGAAAGAGGTGCCCCTGGTGTGCCAAGGAGCC
                                                                                                                         999gcaaaaggctccacaggccccaagggagaaagaggtgcccctggtgtgcaaggagcc 556
                                                                                                                                                                                                                                                                                GACCANAGGGAGAACGTGGACTAAGT-----
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                                                                                                                                                                                                                                                                                                                             TGCAGGGCCTACAGGTCCAGTTGGACCCAAAGGAGAATGGCTCTGCTGGCGAACCTG
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                                                                                                                                                                                                                                                                                                                                                                                 AAGGTCCACGGGGTGAGAAGGGTGATCCAGG-TTGCCAGGACCTATGGGGCTCTCAGGGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            http://image.llnl.gov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="tumor, biopsy sample"
/dev_stage="3 months, virgin"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:10090"
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/clone_lib="NCI_CGAP_Mam1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 428; DB 141;
Pred. No. 6.5e-109;
0: Mismatches 40;
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110

93 aaatgaagagcctctcgcagagatcagtacccaacacctgcaccctagtcatgtgtagcc

AAATGAAGACCTACTCCCACAGAACAACGCCCAGTGCTTGCACCCTGGTCATGTGTAGCT

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ORIGIN
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AUTHORS
TITLE
JOURNAL
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VERSION
KEYWORDS
SOURCE
ORGANISM
                                                                                                                Query Match
Best Local
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                                                                                            Best Local Similarity Matches 487; Conserv
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                     tgctgccctttctctccatgcttgtcttgcttgtacagcccctgggaaatctgggagcag
TGCTCTTCCTCTCTCTGCACTGGTCCTGCTCACACACCCCTGGGCTACCTGGAAGCAG 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov

Tissue Procurement: CLONTECH Laboratories, Inc.
cDNA Library Preparation: CLONTECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
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Plate: LLCM1371 row: c column:
High quality sequence stop: 655.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mar
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 665)
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602502918F1 NIH_MGC_77
mRNA sequence.
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EST.
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                                                                                                                                                                                                        149
                                                                                                                                                                        /organism="Homo sapiens"
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/db_xref="taxon:9606"
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/note="Organ: lung; vector: pDNR-LLB (Clontech); Site_1:
Sfil (ggcgctcggc); Site_2: Sfil (ggccattatggcc); 5' and
3' adaptors were used in cloning as follows: 5' adaptor
sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence:
C' or G and N = A' C' G' Or T'). Average insert size 1.9
kb (range 0.5-4.0 kb). 12/15 colonies contained inserts
by PCR. This library was enriched for full-length clones
and was constructed by Clontech Laboratories (Palo Alto,
CA). Note: this is a NH_MGC Library."
180 c 241 g 94 t lothers
                                                                                        Conservative
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                    24.48; 79.18;
                                                                                      0;
                                                                                                  Score 378.6; DB :
Pred. No. 4.3e-95
                                                                               Mismatches 125;
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650 CCGGGATTGAAGGGGG 665
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                          agcagcaggacctgcc--ggacctgccggtccacagggagctccaggttccagggggccc 628
                                                       ccaggactcaaggggg 644
                                                                                    agaagggtgatccaggtttgccaggacctatggggctctcagggttgcagggccctacag 272
                                                                                                                                                                          GCCCAGTTGGGCCCAAAGGGGACAATGGCTCTGNTGGAGAACCTGGACCAAAGGGAGACA 349
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Search completed: August $\,$ 8, 2001, 05:06:21 $\,$ Job time: $\,$ 7975 $\,$ sec

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